GenCore version 5.1.4\_p5\_4578 Copyright (c) 1993 - 2003 Compugen Ltd.

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Minimum
Maximum
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Maximum Match 100%
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-DB-Tssued_Patents_Na -QFMT=fastap -SUFFIX=rni -MINMATCH=0.1 -LOOPCL=0
-LOOPEXT=0 -UNITS-bits -START=1 -END=-1 -MATRIX-blosum62 -TRANS-human40.cdi
-LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15
-MODE-LOCAL -QUTFMT=pto -NORM-ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=200000000
-USER=US0993234_eCGN_1_1_116_erunat_2703203_115457_15378 -NCPU=6 -ICPU=3
-NO.XLPXY -NO.MAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120
-NARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPDEXT=0.5 -FGAPOP=6 -FGAPEXT=7
-NARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPDEXT=0.5 -FGAPOP=6 -FGAPEXT=7
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-MODEL-frame+_p2n.model -DEV=x1h
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length: 2000000000
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/cgn2_6/ptodata/1/lna/backfiles1.seq:*
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US-08-928-069-2
US-08-928-683A-2
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## ALIGNMENTS

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                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                   PRIOR APPLICATION NATA:
APPLICATION NUMBER: NO. 6153402 Y.
FILING DATE: 06-FEB-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/028,711
FILING DATE: 17-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Yu, Guo-Liang
APPLICANT: Ni, Jian
APPLICANT: Dixit, Vishva
APPLICANT: Gentz, Reiner L.
                      PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Dillon, Patrick J. TITLE OF INVENTION: Death Don
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                       STREET: LIVU New CITY: Washington
                                                                                                                                                                             APPLICATION NUMBER: US/08/815,469 FILING DATE: HEREWITH
                                                                                                                                                            CLASSIFICATION:
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
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5. 6153402
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  US 60/013,285
                                                                                                                     6153402 Yet Assigned
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Sequence 11, Application US
Patent NO. 6462176
GENERAL INFORMATION:
APPLICANT: Ashkenazi, A
TITLE OF INVENTION: Apo
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US-08-928-069-11
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Best Local Similarity:
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TELEPHONE: 202-371-2600
TELEFAX: 202-371-2540
INFORMATION FOR SEQ ID NO: 3:
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FEATURE:
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ATTORNEY/AGENT INFORMATION:
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STRANDEDNESS: double
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                                                                                                                GlySerCysProGluArgCysAlaAlaValCysGlyTrpArg
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                                                                                                                                               TGCCTGCCTGGCTTCTATGAACATGGCGATGGCTGCGTGTCCTGCCCCACGAGCACCCTG
                                                                                                                                                             CysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysProThrSerThrLeu 160
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Matches:
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Best Local Similarity:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
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NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/928,069
FILING DATE: 11-Sep-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/026943
FILING DATE: 09/23/1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compation operating SYSTEM: PC-DO SOFTWARE: WinPatin (Gereur Application Data:
              521
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy
COMPUTER: IBM PC compatible
                                     121
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                                                                                                                                   81
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 1634 base pairs TYPE: Nucleic Acid STRANDEDNESS: Single TOPOLOGY: Linear
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                                                                                                                                                                                                                                                            Percent Similarity:
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Patent No. 6469144
GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: P1007P1
TELECOMUNICATION INFORMATION:
TELEPHONE: 650/952-9881
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 1634 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     581
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 i
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NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible operating SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech) CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Apo-2 LI AND Apo-3 POLYPEPTIDES NUMBER OF SEQUENCES: 28
                                                                                                                                                                                                                                                                                                                                                       SEQUENCE DESCRIPTION: SEQ
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CysGlyAsnSerThrCysLeuValCysProGlnAspThrPheLeuAlaTrpGluAsnHis 60
                                                                LeuPheCysCysArgGlyCysProAlaGlyHisTyrLeuLysAlaProCysThrGluPro 40
                                                                                                                              GlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHisLysLysIleGly
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                                                CTGTTTTGTTGCAGAGGCTGCCCAGCGGGGCACTACCTGAAGGCCCCTTGCACGGAGCCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/828,683A FILING DATE: 31-Mar-1997 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 08/625328
FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: South San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                       TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 1 DNA Way
                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: Linear
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Gaps:
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APPLICANT: Yu, Guo-I
APPLICANT: N1, Jian
APPLICANT: Dixit, Vi
             TELEPHONE: 202-371-260
TELEFAX: 202-371-2540
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                             REFERENCE/DOCKET NUMBER: 14
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
                                                                                                                                         FILING DATE: 17-OCT-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/013,285
FILING DATE: 12-MAR-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: NO. 1
FILING DATE: 06-FEB-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 6
                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
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APPLICANT: Gentz, Reiner L.
APPLICANT: Dillon, Patrick J.
TITLE OF INVENTION: Death Domain Containing Receptors
NUMBER OF SEQUENCES: 17
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatil
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                                                                                                             REGISTRATION NUMBER:
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                                                                                                                               NAME:
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1100 New York Ave.,
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NW, Suite 600
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RESULT 5
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DB:
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; LOCATION:
US-08-815-469-1
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                                                                                                                             Sequence 5, Application US/08928069
Patent No. 6462176
GENERAL INFORMATION:
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44
COMPUTER: IBM PC compatible
                                                                                        CORRESPONDENCE ADDRESS
                                                                                                 TITLE OF INVENTION: Apo-3 NUMBER OF SEQUENCES: 15
                                                                                                                    APPLICANT: Ashkenazi, Avi
                                                 ADDRESSEE:
STREET: 1 |
CITY: Sout|
STATE: Cal:
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FEATURE:
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                                                                           Genentech, Inc.
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RESULT 6
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DB:
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/0
FILING DATE: 09/23/1996
ATTORNEY/AGENT INFORMATION:
NAME: Marschang, Diane L
                                                                                                                                                                    749
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CURRENT APPLICATION DATA:
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STRANDEDNESS: Single
TOPOLOGY: Linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: FILING DATE: 11-Sep CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                       1 GlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHisLysLysIleGly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM:
                                                   GlySerCysProGluArgCysAlaAlaValCysGlyTrpArg
                                                                                           CysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysProThrSerThrLeu 160
                                                                                                                       CysGlnValSerGlnCysValSerSerSerProPheTyrCysGlnProCysLeuAspCys 120
                                                                                                                                                                                                         CTGTTTTGTTGCAGAGGCTGCCCAGCGGGGCACTACCTGAAGGCCCCTTGCACGGAGCCC
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 Application
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11-Sep-1997
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Indels:
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Conservative:
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Best Local Similarity:
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GENERAL INFORMATION:
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101 CysGlnValSerGlnCysValSerSerSerProPheTyrCysGlnProCysLeuAspCys 120
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FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy
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TITLE OF INVENTION: Apo-2 LI AND Apo-3 POLYPEPTIDES
NUMBER OF SEQUENCES: 28
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                                                 GluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGlyTrpPheValGlu 100
                                                                                                                 HisasnSerGluCysalaargCysGlnalaCysaspGluGlnalaSerGlnValalaLeu
                                                                                                                                                                               CysGlyAsnSerThrCysLeuValCysProGlnAspThrPheLeuAlaTrpGluAsnHis 60
                                                                                                                                                                                                                                                                                                 CAGGGCGGCACTCGTAGCCCCAGGTGTGACTGTGCCGGTGACTTCCACAAGAAGATTGGT 508
                                GAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGTGGCTGTAAGCCAGGCTGGTTTGTGGAG 748
                                                                                                CATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGCCTCCCAGGTGGCGCTG
                                                                                                                                                               TGCGGCAACTCCACCTGCCTTGTGTGTCCCCAAGACACCTTCTTGGCCTGGGAGAACCAC
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TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/828,683A FILING DATE: 31-Mar-1997
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Matches:
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Best Local Similarity:
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US-08-928-069-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: P1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 09/23/1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy
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22 PheCysCysArgGlyCysProAla-GlyHisTyrLeuLysAlaProCysThrGluProCy 41
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                                                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                        TYPE:
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                                                         2 GlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHisLysLysIleGlyLeu 21
                                           GGCGGCACTCGTAGCCCCAGGTGTGACTGTGCCGGTGACTTCCACAAGAAGATTGGTCTG
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EDNESS: Single
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Matches:
Conservative:
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Percent Similarity:
Best Local Similarity:
Query Match:
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                                                                                                                     ; SEQUENCE DESCRIPTION: SEQ ID NO: US-08-828-683A-2
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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 433 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: P1007P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
ATTORNEY/AGENT INFORMATION:
NAME: MAISCHANG, Diane L.
REGISTRATION NUMBER: 35,600
DEFERENCE ACCESS MINGELS.
                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/828,683A
FILING DATE: 31-Mar-1997
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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ADDRESSEE: Genentech,
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TITLE OF INVENTION: Apo-2 LI AND
NUMBER OF SEQUENCES: 28
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                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 08/625328
FILING DATE: 1-Apr-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: WinPatin (Genentech)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: South San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET:
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92.03%
64.16%
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               Length:
Matches:
Conservative:
Mismatches:
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                                       TELEFAX: (202) 737-35
INFORMATION FOR SEQ ID NO:
                                                                      NAME: Browdy, Roger L.
REGISTATION NUMBER: 25,618
REFERENCE/DOCKET NUMBER: CA.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 628-5197
                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                      CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Jameson, Bradford A.
APPLICANT: Chappel, Scott C.
TITLE OF INVENTION: HYBRID PROTEINS
NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
                                                                                                                                                          CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Washington
STATE: D.C.
COUNTRY: USA
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          LENGTH:
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        1049 base pairs
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Best Local Similarity:
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                                                                                                                                                                                                                                           Sequence 1, Application US/08910991 Patent No. 6194177
                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Campbe
                                                                                                                          APPLICANT: Jameson, Bradford A.
APPLICANT: Chappel, Scott C.
TITLE OF INVENTION: HYBRID PROTEINS
NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
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                     ZIP: 22207
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                                                                                               E: BROWDY AND NEIMARK
419 Seventh Street N.W., Ste.
                                                                                                                                                                                                             Campbell, Robert K.
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NAME/KEY:
LOCATION:
US-08-910-991-1
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Query Match:
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 37,971
REFERENCE/DOCKET NUMBER: CAN
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 628-5197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 20 February 1996
ATTORNEY/AGENT INFORMATION:
NAME: YUN, Allen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 08/804,166
FILING DATE: 20 February 1997
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA: APPLICATION NUMBER: US
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716
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC
OPERATING SYSTEM:
SOFTWARE: PatentI
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STRANDEDNESS: sing
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                CysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysProThr 157
                                                                   CTCAAT - - - GGGACCGTGCAC - - - - -
                                                                                     LeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThrAsp 137
                                                                                                                                                                   Cys-----GlnValSerGlnCysValSerSerSerProPheTyrCysGlnProCys 117
                                                                                                                                                                                                                                                                                                          HisAsnSerGluCysAlaArgCysGlnAlaCysAspGluGlnAlaSerGlnValAlaLeu
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                                                                                                                                    TATTGGAGTGAAAACCTTTTCCAGTGC------TTCAATTGCAGCCTCTGC
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SYSTEM: PC-DOS/MS-DOS
PatentIn Release #1.0,
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                       Local Similarity:
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STRANDEDNESS: sing
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HisHisAsnSerGluCysAlaArgCysGlnAlaCysAspGluGlnAlaSerGlnValala 79
                                                                                                                                  GlyLeuPheCysCysArgGlyCysProAlaGlyHisTyrLeuLysAlaProCysThrGlu 39
                                              CCAGGGCGGGATACAGTCTGCAGGGAGTGTGAAAAGGGCACCTTTACGGCTTCCCAGAAT 377
                                                                        ProCysGlyAsnSerThrCysLeuValCysProGlnAspThrPheLeuAlaTrpGluAsn 59
                                                                                                                                                                    GGTGACCGGGAGAAGAGGGATAGCTTGTGTCCCCAAGGAAAGTATGTCCATTCTAAGAAC
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Alignment Scores: Pred. No.:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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                                                                                                                      MOLECULE TYPE:
                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
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                                                                                                        FEATURE:
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                                                                                                                                                                                                                                                                                              NAME: Browdy, Roger L. REGISTRATION NUMBER: 25, REFERENCE/DOCKET NUMBER:
                                                                                                                                          TOPOLOGY:
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                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Campbell, Robert K.
APPLICANT: Jameson, Bradford A.
APPLICANT: Chappel, Scott C.
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/804,166
FILING DATE: 20 February 1997
                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
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                                                                                                                                                                                                         COUNTRY:
ZIP: 222
                                                                                                                                                                                                                                  STREET: 419 Sever CITY: Washington STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               81 GluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGlyTrpPheValGlu :::|||::: |||||| |||||||::: :::
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   41 CysGlyAsnSerThrCysLeuValCysProGlnAspThrPheLeuAlaTrpGluAsnHis
                                                     CLASSIFICATION:
                                                                   FILING DATE:
                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                          ADDRESSEE:
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                                                                                                                                                                                           USA
22207
                                                                                                                                                                                                                                                                         419 Seventh Street N.W.,
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                                                                                     us/08/910,991
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Indels:
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Conservative:
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                                               RESULT 14
US-08-050-319B-47
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Best Local Similarity:
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US-08-910-991-7
            Sequence 47, Application US/08050319B Patent No. 5633145
GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (202) 737-35
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/0:
FILING DATE: 20 February
ATTORNEY/AGENT INFORMATION:
NAME: YUN, Allen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 1301 base pairs
TYPE: nucleic acid
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REFERENCE/DOCKET NUMBER: CA
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                          CysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysProThr 157
                                                                                                                                                                                                                                                                                                                                                                                                    GluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGlyTrpPheValGlu :::|||:::
                                                                                                                                                                                                                                                                                                             TATTGGAGTGAAAACCTTTTCCAGTGC------TTCAATTGCAGCCTCTGC
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                                                                                                                                    SerThrLeuGlySerCysProGlu-----ArgCysAlaAlaValCys 171
                                                                                                                                                                   GTGTGCACCTGCCATGCAGGTTTCTTTCTAAGAGAAAACGAGTGTGTCTCCTGT
                                                                                                                                                                                                                                          CTCAAT---GGGACCGTGCAC-----
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Matches:
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Mismatches:
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Best Local Similarity:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 51
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 617-8899
TELEPAX: (415) 327-3231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
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400
                               100 GluCys-----GlnValSerGlnCysValSerSerSerProPheTyrCysGlnPro 116
                                                                                                                                                                                                                                                                                       163 AATTCGATTTGCTGTACCAAGTGCCACAAAGGAACCTACTTGTACAATGACTGTCCAGGC 22
                                                                                                                                                                                                                                                                                                                                                          103 GGGGACAGGGAGAGAGAGATAGTGTGTGTCCCACAGGAAAATATATCCACCCTCAAAAT 162
                                                                                                                                         283 CACCTCAGA---CACTGCCTCAGCTGCTCCAAATGCCGAAAGGAAATGGGTCAGGTGGAG 339
                                                                                                                                                                                                                223
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LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/0 FILING DATE: 10-May-1993 CLASSIFICATION: 435
                                                                                         80 LeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGlyTrpPheVal 99
                                                                                                                                                                                                                                                                                                                        20
                                                                                                                                                                           60 HisHisAsnSerGluCysAlaArgCysGlnAlaCysAspGluGlnAlaSerGlnValAla 79
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 33,208
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CATTATTGGAGTGAAAACCTTTTCCAGTGC-
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: California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    600 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               635 Bryant Street
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M.J.C. Turner, F.M Brennan
VERMION: Modified human TNFalpha (Tumor
VERMION: Necrosis Factor alpha) Receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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                                             US-09-993-234-6_COPY_25_198 (1-174) x US-08-465-982-47 (1-600)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: M.Feldmann, P.W. Gray,
APPLICANT: M.J.C. Turner, F.M Brennan
TITLE OF INVENTION: Modified human TNFalpha (Tumor
TITLE OF INVENTION: Necrosis Factor alpha) Recepto:
                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (415) 327-32 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                             MOLECULE TYPE:
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MEDIUM TYPE: Floppy disk
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Search completed: April 6, 2003, 23:29:46 Job time: 35.3732 secs

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Minimum DB :
Maximum DB :
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-O-/cgn2_1/USPTo_spool/US09993234/runat_27032003_115455_15349/app_query.fasta_1.2346
-DB-GenEmbl -OFMT-fastap -SUFFIX=-ge -MINMATCH-0.1 -LOOPCL-0 -LOOPEXT-0
-UNITS-bits -START-1 -END--1 -MATRIX-blosum62 -TRANS-human40.cdi -LIST-45
-DOCALIGN-200 -THR_SCORE-pct -THR_MAX-100 -THR_MIN-0 -ALIGN-15 -MODE-LOCAL
-OUTFMT-pto -NORM-ext -HEAPSIZE-500 -MINLEN-0 -MAXLEN-2000000000
-USER-US0993234 eCGN 1_1_8534_erunat_27032003_115455_15349 -NCPU-6 -ICPU-3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES-0 -WAIT -LONGLOG -DEV_TIMEOUT-120
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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Yu,G.-L., Ni,J., Gentz,R.L. 44., 2012, R.L. 44., 2012, R.L. 44., 2012, R. 4007-2000;

Patent: US 6153402-A 3 28-MOV-2000;

Location/Qualifiers

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                                                                                                                                    Homo sapiens:

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; I

Mammalia; Eutheria; Primates; Catarrhin; Hominidae;

1 (bases 1 to 1254)

Chinnalyan, A. M., O'Rourke, K., Yu, G. L., Lyons, R. H., Ga

Duan, D. R., Xing, L., Gentz, R., Ni, J. and Dixit, V. M.

Signal transduction by DR3, a death domain-containing

related to TMFR-1 and CD95

Science 274 (5289), 990-992 (1996)

97081063
                                                                   Submitted (29-SEP-1996) Pathology, Un. School, 1301 Catherine St., Box 0602,
                                                                                      2 (bases 1 to 1254)
Chinnaiyan,A.M., O'Rourke,K., Yu,G.-L., Lyons,R.
Duan,R., Xing,L., Gentz,R., Ni,J. and Dixit,V.M.
Direct Submission
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                                                          PheTrpValGlnValLeuLeuAlaGlyLeuValValProLeuLeuLeuGlyAlaThrLeu
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LYDNVDAVPARWKETVTAGLREAEIEAVEVEIGRFRDQQYEMLKRWRQQQPAGLGA
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Goodwin,R.G.
AIR, A Novel Member of Apoptosis
                                                                                                                                                        Direct Submission
Submitted (12-NOV-1996) Biochemistry,
University St., Seattle, WA 98101, USA
                                                                                                                                                                                                                                                             Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae
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e Presente	FEATURES CDS	REFERENCE AUTHORS TITLE JOURNAL	REFERENCE AUTHORS TITLE JOURNAL MEDLINE	RESULT 4 HSU94501 LOCUS DEFINITION ACCESSION VERSION KEYWORDS SOURCE ORGANISM	u u .	Qy       3         Db       9         Qy       3         Db       10         Qy       3         Db       10	Qy 2 Db 8 Qy 3 Db 9
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/function="mediates apoptosis" /note="nARD-la; membrane protein; similar to Fas and TNEF-R1; contains a death domain" /codon_start=1 /product="lymphocyte associated receptor of death la" /protein_id="AAC51306.1" /db_xref="G1:2071949" /translation="MEQRERGCAAVAAALLLVILIGARAQGGTRSBRCDCAGDEHKKIG LFCCRGCPAGHYLKAPCTEPCGNSTCLVCPQDTFLAWENHHNSECARCQACDEQASQV ALENCSAVADTRCGCKPGWFVECQVSQCYSSSPFYCQPCLDCGALHRHTELLCSRRDT DCGTCLPGFYEHGDGCVSCPTSTLGSCPERCAAVCGWRQMFWVQVLLAGLVVPLLLGA	Location/Qualifiers 11254 /organism="Homo sapiens" /db_xref="taxon:9606" /cell_line="HeLa" /cell_type="lymphocyte" 11254		Mammalia; Eutheria; Primates; Cranlata; Vertebrata; Eutheleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  1 (bases 1 to 1254) Screaton,G.R., Xu,X.N., Olsen,A.L., Cowper,A.E., Tan,R., McMichael,A.J. and Bell,J.I. LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)	1254 bp mRNA hocyte associated receptor of 31:2071948	ATCOMPACTO TOGAGETIGAGATCGGCCGCTTCCGAGACCAGCAGTACGAGATGCTCAAG 1140 ATGTTPATGGINGINGTNFTOALAGIYLEUGIYALAVALTYFALAALALEUGIUATGMEt 400	ProAlaGlySerProAlaMetMetLeuGlnProGlyProGlnLeuTyrAspValMetAsp 340	TrpThrProGlyTyrProGluThrGlnGluAlaLeuCysProGlnValThrTrpSerTrp 300

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2 (bases 1 to 1634)
Marsters, S.A., Sheridan, J.P.,
Goddard, A.D., Bauer, K.D. and
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NF-kB; TNF receptor family me
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Ashkenazi,A.J., Goddard,A., Godowski,P.J., Gu Hillan,K.J., Marsters,S.A., Pan,J., Pitti,R.M Stone,D.M., Watanabe,C.K. and Gwod,W.I. Compositions and methods for the treatment of Patent: WO 0153486-A 23 26-JUL-2001; Genentech, Inc. (US)
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1 (bases 1 to 1257)
Screaton,G.R., Xu,X.N., Olsen,A.L., Cowper,A.E., Tan,R., McMichael,A.J. and Bell,J.I.
LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing
Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
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                                                                 TGCCTAGACTGCGGGGCCCTGCACCGCCACACGCCTACTCTGTTCCCCGCAGAGATACT
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/product="lymphocyte associated receptor of death 1b"
/product="lamac51307.1"
/product="lymphocyte associated receptor of death 1b"
/protein_id="AGC51307.1"
/product="lymphocyte associated receptor of death 1b"
/product="lymphocyte"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note="insertion compared to LARD-1a, deposited in GenBar
Accession Number U94501, probably represents alternative
3' splice site"
a 421 c 408 g 226 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note="LARD-1b; membrane
TNF-R1; contains a death
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/db_xref="taxon:9606"
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Chaudhary, P.M. and Hood, L.E.
Direct Submission
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                                                                                     Submitted
                                                                                                                     Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                Homo sapiens.
                                                                             Washington,
                                                                                                                                                                                          HSU83597
                                                                                                                                       sapiens
                                                                                                                                                                                 death domain
        /organism="Homo sapiens"
/db_xref="taxon:9606"
/chromosome="1"
/map="1p36"
1 . 1528
                                                                           (03-JAN-1997)
n, 1705 NE Paci
/gene="DDR3"
                                                          Location/Qualifiers
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                                                                           997) Molecular
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3 (DDR3) mRNA,
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                                                                           Biotechnology, University of t, HSB-K360, Seattle, WA 98195
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/gene="DDR3"
/gene="DDR3"
/function="mediates apoptosis when cross-linked"
/function="mediates apoptosis when cross-link
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/protein_id="ABAB41432.1"
/protein_id="ABAB41432.1"
/db_xref="GI:1800293"
/db_xref="GCAAVAAALLLVLLGARAQGGTRSPRCDCAGDPHKKIGLFCCRG
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AVADTRGGCKPGWFVECQVSGCVSSSPFYCQPCLDCGALHRHTLACSRRDTDCGTCL
LGFYEHGDGCVSCPTSTLGSCPERCAAVCGWAMFWVQVLLAGLVVPLLLGATLTNTYT
RHCWPHKPLVTADEAGMEALTPPPATHLSPLDSAHTLLAPPDSSEKICTVQLVGNSTT
PGYPETGEALCPQVTWSWDQLPSRALGPAAAPTLSPESPAGSPAMMLQPGPQLYDVMD
AVPARRWKEFVRTLGLREAEIEAVEVEIGRFRDQQYEMLKRWRQQQPAGLGAVYAALE
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                                     Direct Submission
Submitted (18-0CT-1996) J
Tausanne, 155 Chemin des
                                                                                                                                       Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Bukaryota; Metazoa; Chordata; Craniata; Hominidae; Homo.

1 (bases 1 to 1557)

1 (bases 1, to 1557)

Bodmer, J.L., Burns, K., Schneider, P., Hofmann, K., Steiner, V.,

Thome, M., Bornand, T., Hahne, M., Schroter, M., Becker, K., Wilson, A.,

French, L.E., Browning, J.L., MacDonald, R. and Tschopp, J.

TRAMP, a novel apoptosis-mediating receptor with sequence homology to tumor necrosis factor receptor 1 and Fas(Apo-1/CD95)

Immunity 6 (1), 79-88 (1997)
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2 (base
                                                                         2 (bases 1 to 1557)

Bodmer, J.L., Burns, K., Schneider, P., Hofmann, K., Steiner
Thome, M., Bornand, T., Hahne, M., Schroter, M., Wilson, A.,
French, L.E., Browning, J.L., MacDonald, R. and Tschopp, J.
                                  Lausanne, 1:
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 /organism="Homo
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                                                      University
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VMDAVPARRWKEFVRTLGLREAEIEAVEVEIGRFRDQQYEMLKRWRQQQPAGLGAVYA
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86 a 508 c 477 g 286 t
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 1355)
Screaton,G.R., Xu,X.N., Olsen,A.L., Cowper,A.E., Tan,R.,
McMichael,A.J. and Bell,J.I.
LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing
Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
                                 Direct Submission
Submitted (19-MAR-1997) N
Molecular Medicine, John
                                                                 2 (bases 1 to 1355)
Screaton, G.R.
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spliced, complete cd:
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Human lymphocyte associated receptor
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                                   Molecular Immunology Group, Institute
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/note="LARD-2; similar to Fas
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/protein_id="AAC51308
/db_xref="GI:2071953"
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Sequence 1 from p
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Yu,G.-L., N1,J., Gentz,R.L. and Dillon,P.J.
Death domain containing receptors
Patent: US 6153402-A 1 28-NOV-2000;
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red. No.:  2367-00  Scroen: Similarity:  97.394  Scroen: Similarity:  97.394  Missacches:  80-993-234-6 (1-417) x ARI19656 (1-1783)  11111111111111111111111111111111111				•
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Patent:
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kitson, J.
Direct Submission
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Pun,K.T., Grinham,C.J., Brown,R. and Farrow,S.N.
A death-domain-containing receptor that mediates
Nature 384 (6607), 372-375 (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                H.sapiens mRNA for WSL-LR, WSL-S1 and WSL-S2
Y09392
Y09392.1 GI:1669690
wsl-1 gene; WSL-1R protein; WSL-S1 protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 1743)
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LYDVMDAVPARRWKEFVRTLGLREAEIEAVEVEIGLFRDQYEMLKHWRQQQPAGLGA
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                                                                                                                                                                                                         join(69. .610,768. .882)
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1. .1742
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Percent Similarity:
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Query Match:
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               ThrSerThrLeuGlySerCysProGluArgCysAlaAlaValCysGlyTrpArg-----
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                                                                                      AspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysPro 180
                                                                                                                                                       CysLeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThr
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TGCCTAGACTGCGGGGGCCCTGCACCGCCACACGGCTACTCTGTTCCCGCAGAGATACT
                                                                                                                                                                                                      TGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGCCAACCA
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/number=3
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DCGTCLPGFYEHGDGCVSCPTSTLGSCPERCAAVCGWRQSRWCAGNARGRTGMDRGEA
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GlnMetPheTrpValGlnValLeu

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JOURNAL
MEDLINE
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AUTHORS
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SOURCE
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                                                                                                                                                                              Homo sapiens AF026070 AF026070.1 G
                                                                                                     Salles, G.
A new deat
Direct Submission
Submitted (19-SEP-1997) Hematology, CTRE Hospitaller
Chemin du Grand Revoyet, Pierre Benite 69495, France
Location/Qualifiers
1. 1669
                                                                          lines and non-Hodgkin's lymphomas
Blochem. Blophys. Res. Commun. 242 (2), 376
98113360
                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 1669)
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Warzocha, K., Ribeiro, P.,
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Percent Similarity:
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QQQPAGLGAVYAALERWGLDGCVEDLRSRLQRGP"
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/note="Apo-3;
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spliced product;
extracellular don
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patients with non-Hodgkin's lymphoma"
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ECYSThrValGlnLeuValGlyAsnSerTrpThrProGlyTyrProGluThrGlnGluAl 291
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Search completed: April 6, 2003, 21:07:27 Job time: 4876.17 secs

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Query

SUMMARIES

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-MODEL-frame+_p2n.model -DEV-xlh
-Q-/cgn2_1/USPTO_spool/US09993234/runat_27032003_115454_15312/app_query.fasta_1.
-DB-N_Geneseq_101002 -QFMT=fastap -SUFFIX=rng -MINMATCH=0.1 -LOOPCL=0
-LOOPEXT=0 -UNITS-bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi
-LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15
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-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPDP=6 -FGAPEXT=7
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Copyright (c) 1993 - 2003
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Pred. No. is the number of results predicted by chance to have score greater than or equal to the score of the result being pland is derived by analysis of the total score distribution. printed, RESULT 1
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Result No. a Q 2267 2254.5 2208 1556.5 1556.5 1404 1051 1039.5 394.5 .378.5 Score 1000.0 1100.0 10 Match Length 14825 10797 1251 1438 1438 186 2130 2173 1368 1368 1368 2062 2088 2088 2111 2111 2111 2111 2111 2175 2175 1254 1254 1634 1634 1634 1662 1847 1783 1783 1783 1783 1743 2141 511 511 511 1334 511 511 1704 멂 片 ABK63694 AAQ406284 AAQ49932 AAA95105 AAQ20973 AAQ24440 AAQ10883 AAC83946 AAZ09170 AAH48859 ABK844039 ABK42690 AAV28701 AAT91179 AAL47187 ABL67258 ABK40265 AAV28700 AAT89426 AAX00924 AAC68776 ABL64119 AAF83770 AAL47185 AAL47186 AAC91477 AAQ50870 AAQ10955 AAQ12215 AAT89427 AAX00925 AAC68777 AAQ06285 ABQ47530 ABK13194 AAQ90513 ABN95862 AAZ48475 Human rheumatoid a Human PRO779 cDNA. cDNA encoding huma Human apoptosis in Death domain conta Death domain conta Human death domain Breast cancer rela Type I TNF recepto Human Tumour Necro Oligonucleotide fo Oligonucleotide fo Human TNFR1 coding
TNF-alpha binding
Encodes TNF-alpha
30kD TNF inhibitor
Human 30 kDa TNF i Human tumour necro Human TNFBP-associ Human cDNA differe Gene #2360 used to Mouse apoptosis in Human apoptosis pr Human rheumatoid a Thyroid cancer rel Description Oligonucleotide Oligonucleotide p55 TNF-R gene. H p55 Tumour necrosi Encodes human 55kD Rat sequence diffe Rat Tumour Necrosi Lambda-derived TNF Nucleotide sequenc Human DR3 gene ass Genomic sequence # Oligonucleotide fo Plasmid Tumour Nec Human tumour necro death domain apoptosis pr genomic DNA domain conta domain conta diffe

ALIGNMENTS

AAT89427 standard; cDNA; 1254 BP.

02-MAR-1998 (first entry)

AAT89427;

Death domain containing receptor DR3 cDNA.

Death domain containing receptor; DR3; human; apoptosis; inflammation; NF-kappaB; ds. Death

Homo sapiens

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                                                                                                                                                                                                                                                                                                                     Alignment
                                                                                                                                                                                                                                                                                                                                                       This cDNA clone codes for human death domain containing receptor CDR3 (see AAW31517), a novel member of the tumour necrosis factor receptor family. It was isolated from a HUVEC cDNA library. Related death domain containing receptor DR3-V1 cDNA (see AAR39426) was isolated from a human testis tumour cDNA library. The genes consisted the second dentified in cDNA libraries of foetal liver, foetal brain, tonsil and leukocyte. Nucleic acids encoding full-length or mature DR3, or the extracellular, transmembrane, intranslelular or especially the death domain of DR3, can be used to produce recombinant polypeptides in transformed host cells. These polypeptides can be used to treat diseases and disorders associated with increased apoptosis and for treating inflammatory diseases and disorders.
                                                                                                                                                                                                                                                                                                          No.:
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                    GlnValAlaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGly
                                                            GGGGCCCGGGCCCAGGGCACTCGTAGCCCCAGGTGTGACTGTGCCGGTGACTTCCAC
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Death domain

containing receptor polypeptide (DR3)

encoding

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12-MAR-1996;
17-OCT-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Treating graft-versus-host disease, cancer, immunodeficiency or an autoimmune disease comprising administering an antibody to Death Decontaining Receptor proteins and a second therapeutic agent -
                                                                                                                                                                                                                                                                                                           Sequence 1254 BP; 201 A; 420 C;
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                                   (GENT/)
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(UNMI) UNIV MICHIGAN.
(YUGG/) YU G.
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28-MAY-1999;
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autoimmune disorder; inflammation; cardiovascular disorder; infection;
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CysThrGluProCysGlyAsnSerThrCysLeuValCysProGlnAspThrPheLeuAla
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) NI J.
) GENTZ R L.
) DILLON P J.
) DIXIT V M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ni J,
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                                                US-09-993-234-6 (1-417) x AAT91180 (1-1634)
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                                                                                                                                                                                                  CDNA clone FH20.57 (ATCC 55820) codes for a novel polypeptide (see W26709), designated Apo-3, that stimulates or induces apoptotic cactivity in mammalian cells. It was isolated from a human foetal cheart cDNA library by screening with probes (see T91183-84) based on an EST sequence (GenBank locus W71984) that showed homology to the intracellular domain of human TNFR1 and CD95. Amino acid cresidues 1-181 of Apo-3 are identical to another novel apoptosis can be used diagnostically for tissue-specific typing and to produce recombinant Apo-3 polypeptides, especially the extracellular domain (amino acids 1-198) or death domain (amino acids 198) and to produce recombinant Apo-3 can be used to induce apoptosis or Cards 338-417). Apo-3 can be used to induce apoptosis or NF-kappa-N- or JNK-mediated gene expression for therapeutic purposes. Non-human transgenic animals containing cells that express Apo-3 nucleic acid, and knockout animals containing cells that have an altered Apo-3 gene, can be used in drug
                                                                                                                                                                                                                                                                                                                                                                                                                                          Polypeptide(s) Apo-3 and Apo-2LI - useful for regulating apoptosis in mammalian cells
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apoptosis protein Apo-3 cDNA clone FH20.57
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IleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGluMetLeuLys 38 	AlaValProAlaArgArgTrpLysGluPheValArgThrLeuGlyLeuArgGluAlaGlu 36	ProAlaGlySerProAlaMetMetLeuGlnProGlyProGlnLeuTyrAspValMetAsp 34	AspGlnLeuProSerArgAlaLeuGlyProAlaAlaAlaProThrLeuSerProGluSer 32 	TrpThrProGlyTyrProGluThrGlnGluAlaLeuCysProGlnValThrTrpSerTrp 30	LeuLeuAlaProProAspSerSerGluLysIleCysThrValGlnLeuValGlyAsnSer 28 	MetGluAlaLeuThrProProProAlaThrHisLeuSerProLeuAspSerAlaHisThr 26	ThrTyrThrTyrArgHisCysTrpProHisLysProLeuValThrAlaAspGluAlaGly 24	PheTrpValGlnValLeuLeuAlaGlyLeuValValProLeuLeuLeuGlyAlaThrLeu 22 	ThrSerThrLeuGlySerCysProGluArgCysAlaAlaValCysGlyTrpArgGlnMet 20	AspCysClyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysPro 18	CysLeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThr 16	TrpPheValGluCysGlnValSerGlnCysValSerSerSerProPheTyrCysGlnPro 14	GlnValAlaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGly 12 	TrpGluAsnHisHisAsnSerGluCysAlaArgCysGlnAlaCysAspGluGlnAlaSer 10 	CysThrGluProCysGlyAsnSerThrCysLeuValCysProGlnAspThrPheLeuAla 80 	LysLysIleGlyLeuPheCysCysArgGlyCysProAlaGlyHisTyrLeuLysAlaPro 60 	GlyalaargalaGlnGlyGlyThrargSerProArgCysaspCysalaGlyAspPheH1s 40 
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GlyAlaArgAlaGlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHis
                                       invention relates to a method of diagnosing chronic rheumatoid ritis by digesting human genomic DNA with ECORI and hybridising a probe containing a fragment of the present sequence which esents DNA encoding a transmembrane protein. The method is used diagnosis of chronic rheumatoid arthritis, and for developing not be a sequence of the present sequence.
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                                                                                                                                                  The present invention relates to the human DR3 gene, which is associated with rheumatoid arthritis. Certain mutations in the gene can be linked to the disease. The sequences can be used to evaluate disease onset and its possibility and to provide therapy and remedies. The present sequence is a coding sequence described in the exemplification of the invention.
                                                                                                                                                                                                               Genomes, particularly DR3 genomic DNA, participating arthritis via mutation, useful in evaluating disease possibility and providing therapy and remedies -
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Y; gene; ss.
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                                                                                          IleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGluMetLeuLys
                                                             GCGGTCCCAGCGCGCGCTGGAAGGAGTTCGTGCGCACGCTGGGGCTGCGCGAGGCAGAG
                                                                    AlaValProAlaArgArgTrpLysGluPheValArgThrLeuGlyLeuArgGluAlaGlu
                                                                                                                                                                                        MetGluAlaLeuThrProProProAlaThrHisLeuSerProLeuAspSerAlaHisThr
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                               ATCGAAGCCGTGGAGGTGGAGATCGGCCGCTTCCGAGACCAGCAGTACGAGATGCTCAAG
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15-MAR-2000;
20-MAR-2000;
21-MAR-2000;
The present sequence is one of thirty three nucleic acids encoding propolypeptides. The PRO polypeptides, anti-PRO antibodies, agonists and
                                                Thirty three nucleic acids encoding PRO polypeptides which are useful in the diagnosis and treatment of immune related disorders, e.g. systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis, thyroiditis and diabetes mellitus -
                               Claim 48; Fig 33; 218pp; English.
                                                                                                    WPI; 2001-025253/03.
P-PSDB; AAB50918.
                                                                                                                                         Ashkenazi AJ, Baker KP, Chan
Hebert C, Henzel W, Kabakoff
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22-FEB-2000;
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20-JUL-1999;
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29-OCT-1999;
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L5-SEP-1999;
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2000WO-US00376
2000WO-US03565
2000WO-US04341
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2000WO-US04914
2000WO-US04914
2000WO-US06884
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2000WO-US07532
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ic; cardiant; antianaemic; immunosuppressive; ant
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2000WO-US14042.
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99US-0146222.
99WO-US20111.
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Shelton DL,
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,, Tumas D, Watanabe CK;
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antagonists are useful for treating and diagnosing immune related disorders such as systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis, juvenile chronic arthritis, spondyloarthropathies, systemic sclerosis, idiopathic inflammatory myopathies, Sjogren's syndrome, systemic vasculitis, sarcoidosis, autoimmune haemolytic anaemia, autoimmune thrombocytopaenia, thyroiditis, diabetes mellitus, immune-mediated renal disease, demyelinating diseases of the central and peripheral nervous systems (such as multiple sclerosis, idiopathic inflammatory demyelinating polyneuropathy), hepatobiliary diseases (such as infectious, autoimmune chronic active hepatitis, primary biliary cirrhosis, granulomatous hepatitis and sclerosing cholangitis), inflammatory bowel disease, gluten-sensitive enteropathy and Whipple's disease, autoimmune or immune-mediated skin diseases (such as bullous skin diseases such as asthma, allergic rhinitis, atopic dermatitis, food hypersensitivity and urticaria), immunological diseases of the lung (such as eosinophilic pneumonias, idiopathic pulmonary fibrosis including graft rejection and graft-versus-host diseases.

Sequence 1662 BP; 302 A; 540 C; 531 G; 289 T; 0 other;

Pred. No Score: Percent Best Loc Query Ma	No.: 7.19e-122 Length: 162 s: 2323.00 Matches: 417 ent Similarity: 100.00% Conservative: 0 Local Similarity: 100.00% Mismatches: 0 r Match: 100.00% Indels: 0 Caps: 0	
us-09	)-993-234-6 (1-417) x AAC91477 (1-1662)	
Qy	1 MetGluGlnArgProArgGlyCysAlaAlaValAlaAlaAlaLeuLeuLeuValLeuLeu 20	
망	103 ATGGAGCAGCGGCGCGGGGCTGCGCGGCGGCGCGCCCCCCCC	
Qy	21 GlyAlaArgAlaGlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHis 40	
DЬ	163 GGGCCCGGGCCCAGGCGCACTCGTAGCCCCCAGGTGTGACTGTGCCGGTGACTTCCAC 222	
Qy	41 LysLysIleGlyLeuPheCysCysArgGlyCysProAlaGlyHisTyrLeuLysAlaPro 60	
Db	223 AAGAAGATTGGTCTGTTTTGTTGCAGAGGCCTGCCCAGCGGGGCACTACCTGAAGGCCCCT 282	
Qy	61 CysThrGluProCysGlyAsnSerThrCysLeuValCysProGlnAspThrPheLeuAla 80	
Db	283 TGCACGGAGCCCTGCGGCAACTCCACCTGCCTTGTGTGTCCCCCAAGACACCCTTCTTGGCC 342	
Qy	81 TrpGluAsnHisHisAsnSerGluCysAlaArgCysGlnAlaCysAspGluGlnAlaSer 100	
DЬ	343 TGGGAGAACCACATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGCCTCC 402	
Qy	101 GlnValAlaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGly 120	
Db	403 CAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCCAGGC 462	
Qy	121 TrpPheValGluCysGlnValSerGlnCysValSerSerSerProPheTyrCysGlnPro 140	
ф	463 TGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGCCAACCA 522	
Qy	141 CysLeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThr 160	
рь	523 TGCCTAGACTGCGGGGCCCTGCACCGCCACACACGGCTACTCTGTTCCCGGCAGAGATACT 582	
Qy	161 AspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysPro 180	
뮹	583 GACTGTGGGACCTGCCTGCCTTCTATGAACATGGCGATGGCTGCGTGTCCTGCCCC 642	
Qy	181 ThrSerThrLeuGlySerCysProGluArgCysAlaAlaValCysGlyTrpArgGlnMet 200	
Db	643 ACGAGCACCCTGGGGAGCTGTCCAGAGCGCTGTGCCGCTGTCTGT	

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RRSULT 8
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08-MAR-1999;
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neuronal disorder;
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99WO-US05028.
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99WO-US12252.
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sorder; stromal disorder; blastocoelic
immune disorder; anglogenic disorder;
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17-AUG-1999
31-AUG-1999
01-SEP-1999
15-SEP-1999
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05-JAN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to the isolation of novel human PRO polypeptides and the polynucleotide sequences encoding them. The PRO polypeptides, agonists, antagonists or anti-PRO antibodies are useful for treating benign or malignant tumours (e.g. renal, kidney, bladder, breast, etc), leukaemias and lymphoid malignancies, other disorders such as neuronal, gllal, astrocytal, hypothalamic, glandular, macrophagal, stromal and blastocoelic disorders, inflammatory, lumune and angiogenic disorders. The polynucleotide sequences are also useful in gene therapy. ABK40254-ABK40288 encode for the human PRO polypeptides of the invention.
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Watanabe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI;
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                                                                                                                                 TrpGluAsnHisHisAsnSerGluCysAlaArgCysGlnAlaCysAspGluGlnAlaSer
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                                                       TGGGAGAACCACCATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGCCTCC
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CAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCCAGGC

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RESULT 9
AAV28700
ID AAV2
XX
AC AAV2
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AC AAV2
XX
DT 20-A
XX
DE Huma
XX
KW Apop
KW Type
Apoptosis inducing receptor; AIR protein; Type I transmembrane protein; tumour cell
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                                                                                                                                                                              AlaValProAlaArgArgTrpLysGluPheValArgThrLeuGlyLeuArgGluAlaGlu
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                   apoptosis inducing receptor coding sequence.
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                                                              standard;
                                   entry)
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 human;
death;
cell death autoimmune
death regulator;
immune disease;
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This sequence encodes the human apoptosis inducing receptor (AIR) of the invention. AIR is a Type I transmembrane protein, soluble forms of which can be used to regulate cell death in a therapeutic setting. Soluble AIR can also be used in vitro to block apptosis or AIR-expressing cells, or to screen agonists or antagonists of AIR activity. The cytoplasmic domain of AIR can be used to develop assays for inhibitors of AIR-induced cell death, which is useful to regulate cell death in a therapeutic setting as well as in vitro. Agonists of AIR activity can be used to kill tumour cells that express AIR, or T cells expressing AIR in autoimmune diseases.
                                                                                                                                                                                                                                                                                                                Claim
                                                                                                                                                                                                                                                                                                                                                              DNA encoding apoptosis inducing receptor - which transmembrane protein, useful for regulating cell
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DB; AAW57045.
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Sequence 1847 BP; 316 A; 605 Ç 579 G; 347 7 0 other;

Alignment Pred. No.: Percent Similarity: Best Local Similarity: . . Match: 8.05e-122 2323.00 100.00% 100.00% 100.00% 100.00% Conservative: Mismatches: Indels: Gaps: Length: Matches: 1847 417 0 0 0

US-09-993-234-6 (1-417) x AAV28700 (1-1847)

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                  domain
containing
                 containing receptor DR3-V1 cDNA.
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receptor;
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DR3-V1; human;
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                                                                                                                                                                                                                                    This cDNA clone, deposited as ATCC 97456, codes for human death domain containing receptor DR3-V1 (see AAW31516), a novel member of the tumour necrosis factor receptor family. It was isolated from a cDNA library derived from cells of a human testis tumour. Related death domain containing receptor DR3 cDNA (see AAW89427) was isolated from a HUVEC cDNA library. The genes have also been identified in cDNA libraries of foetal liver, foetal brain, tonsil and leukocyte. Nucleic acids encoding full-length or mature DR3-V1 can be used to produce recombinant polypeptides in transformed host cells. These polypeptides can be used to treat diseases and disorders associated with horeased apoptosis. Antagonists, such as antibodies raised against DR3-V1, can be used to treat diseases and disorders associated with horeased apoptosis and for treating inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (HUMA-)
                                                                                                                                                                                                                                                                                                                                                                                                                        Death domain containing receptor polypeptide(s) DR3 and DR3-V1 - for activation of apoptosis and NF-kappaB, antagonists can be used to treat inflammatory diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CDS
                                                                                                                                                                                                                                                                                                                                                                                                   Claim 2; Page 71-73; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1997-470812/43.
P-PSDB; AAW31516.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      inflammation;
                                                                                                                                                                                                        Sequence 1783
                                                                                                                                                                                                                              diseases and disorders.
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UNIV MICHIGAN.
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DR3-V1"

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GACGTGATGGACGCGGTCCCAGCGCGCGCGCTGGAAGGAGTTCGTGCGCACGCTGGGGGCTG
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12-MAR-1996;
17-OCT-1996;
                                                                                                                                                                                                                                                                                                                              The invention provides nucleotide sequences encoding death domain containing receptor polypeptides DR3-V1 and DR3. The DR3-V1 cDNA clone is contained in ATCC deposition No. 97456 and the DR3 cDNA clone is contained in ATCC deposition No. 97757. Recombinant vectors comprising the nucleic acid sequences and optionally the leader sequences are used for the recombinant production of the proteins. The present sequence represents a cDNA encoding the death domain containing receptor polypeptide (DR3-V1).
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DB; AAW95537.
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96US-0028711.
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                                                                                                        death domain containing receptors, designated DR3 and DR3-V1. These receptors are involved in apoptosis, and the sequences given can the treatment of cancers, infections, cardiovascular disorders arrhythmias, ischaemia, aneurysms, arterial occlusive diseases, en and congenital heart defects, neurodegenerative diseases including Alzheimer's and parkinson's diseases, autoimmune disease such as sclerosis, arthritis, diabetes, Graves' disease, asthma and psorie
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cancer; colon; breast; ovary; oesophagus; kidney; thyroid; h; lung; prostate; pancreas; carcinoma; antitumour; cancerous; atic; gene therapy; antineoplastic; Wilm's tumour; adenocarcinoma;

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02-OCT-2000; 2000US-237315P.
02-OCT-2000; 2000US-237315P.
03-OCT-2000; 2000US-237598P.
03-OCT-2000; 2000US-237604P.
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01-NOV-2000; 2000US-245084P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Screening for anti-neoplastic agent involves exposing cells to chemical agent to be tested for anti-neoplastic activity, and determining a change in expression of a gene of a signature generation.
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Oy 407 GluAspLeuArgSerArgLeuGlnArgGlyPro 417	387 ProAlaGlyLeuGlyAlaValTyrAlaAlaLeuGluAr 	367 GluIleGlyArgPheArgAspGlnGlnTyrGluMetLeuLy:	Oy 347 TrpLysGluPheValArgThrLeuGlyLeuArgGluAlaGluIleGluAlaValG	Qy 327 MetMetLeuGlnProGlyProGlnLeuTyrAspValMetAspAlaValProAlaAr	Qy 307 AlaLeuGlyProAlaAlaAlaProThrLeuSerProGluSerProAlaAlaNaProThrLeuSerProGluSerProAlaNaProThrLeuSerProAlaNaProThrLeuSer	10 2	Qy 267 SerSerGluLysIleCysThrValGlnLeuValGlyAsnSerTrpThrProGlyTyrPro	Qy 247 ProProAlaThrHisLeuSerProLeuAspSerAlaHisThrLeuLeuAlaProProAsp 	Qy 227 CysTrpProHisLysProLeuValThrAlaAspGluAlaGlyMetGJ 	Qy 207 LeuAlaGlyLeuValValProLeuLeuLeuGlyAlaThrLeuThrT	199	669 AGGTGGTGTGCTGGGAATGCGCGTGGGAGAACTGGGATGGACCG	QУ 198	Qy 181 ThrSerThrLeuGlySerCysProGluArgCysAlaAlaValCysGlyTr	Qy 161 AspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCys\	Qy 141 CysLeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgAr 	Qy 121 TrpPheValGluCysGlnValSerGlnCysValSerSerSerProPheTyrCysGlnP	Qy 101 GlnValAlaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGl 
	gMetGlyLeuAspGlyCysVal 	3ArgTrpArgGlnGlnGln 	eGluAlaValGluVal                 CGAAGCCGTGGAGGTG	aValProAlaArgArg             GTCCCAGCGCGCGC	DAlaGlySerProAla 	pGlnLeuProSerArg              CAGTTGCCCAGCAGA	PThrProGlyTyrPro            accccTGGCTACCCC	uLeuAlaProProAsp 	tGluAlaLeuThrPro 	rTyrThrTyrArgHis 	tPheTrpValGlnValLeu             TTCTGGGTCCAGGTGCTC	AGGGGAGGCGGGTGAG		3GlyTrpArg           GGCTGGAGGCAGAGT	YCYSValSerCysPro 	SSERARGARGASPThr 	OPheTyrCysGlnPro 	3GlyCysLysProGly            GGCTGTAAGCCAGGC
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                                    The invention relates to a composition comprising a biologically active TR3-specific binding agent (I) that binds to TR3 and inhibits the proliferation of cells expressing TR3. (I) identified by the methods are useful for treating a subject suspected of having a disease associated with a proliferation of cells expressing TR3 especially leukemias or 1ymphomas or a T-cell mediated disease especially autoimmune diseases such as myasthenia gravis, systemic lupus erythamatosus, rheumatoid arthritis, diabetes, multiple sclerosis, sarcoidosis, myocarditis, thyroiditis and tumours. (I) is also useful for treating a subject suspected of having graft-versus-host disease, rejection of a transplanted organ such as heart, liver, lung, kidney pancreas, bowel, skin or an appendage, or inflammatory diseases, allergies and contact dermatitis. The present sequence represents the nucleotide sequence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Composition for treatment of T-cell mediated disease e.g. an cancer comprises a biologically active TR3-specific binding especially a monoclonal antibody -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure;
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P-PSDB; AAB84941.
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                AspGlnLeuProSerArgAlaLeuGlyProAlaAlaAlaProThrLeuSerProGluSer
                                                      LeuLeuAlaProProAspSerSerGluLysIleCysThrValGlnLeuValGlyAsnSer
                                                                                            TG-GAGGCTCTGACCCCACCACCGGCCACCCATCTGTCACCCTTGGACAGCGCCCACACC
                                                                                                    MetGluAlaLeuThrProProProAlaThrHisLeuSerProLeuAspSerAlaHisThr
                                                                                                                                            ThrTyrThrTyrArgHisCysTrpProHisLysProLeuValThrAlaAspGluAlaGly
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Human; DR3; rheumatoid arthritis; antirheumatic; antiarthritic;
gene therapy; gene; ds.
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1109..1476
/*tag= g
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974..1108
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636..756
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1645..1742

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757..973
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1..127
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                                      US-09-993-234-6 (1-417) x AAL47185
                                                                                                                       Alignment Scores:
                                                                                                                                                             The present invention relates to the human DR3 gene, which is associated with rheumatoid arthritis. Certain mutations in the gene can be linked to the disease. The sequences can be used to evaluate disease onset and its possibility and to provide therapy and remedies. The present sequence is the gene of the invention.
                                                                                                                                                                                                                                            Genomes, particularly DR3 genomic DNA, participating in rheumatoid arthritis via mutation, useful in evaluating disease onset and its possibility and providing therapy and remedies -
                                                                                                                                                                                                                                                                                                                                                             24-OCT-2000; 2000JP-0324296
27-MAR-2001; 2001JP-0090546
30-MAR-2001; 2001JP-0099990
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585 GAGGCCCCACGTGGGCAGCTGCGCTGAGAGTCAGCAGCACCTGTCCCCCAGGCGCTCCTC
           Konishi
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4022..4202
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4434..4825
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3334..3529
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3125..3225
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RESULT 15 AAL47185

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182 182	Qy		157ArgArgAspThrAspCysGlyThrCysLeuProGl 168	Qy
2742 GCGCCCGGCCTCCATTCAAGTCTTTATTGAATATCTGCTATGTTCTACACACTGTTCTAG 2801	Db		1663 GGCTCTCTACTCCCAGACCCCCTTCTCCCTGCCTGACCCACTCCTGTCCCATGGTGACGC 1722	망
182 182	Qy		156 156	Qy
2682 GACCTCGTGATCCACCCGCCTTGGCCTCCCAAAGTGCTGGGATTACAGGCATGAGCCACC 2741	рь		1604 CCTAGACTGCGGGGCCCTGCACCGCCACACACGGCTACTCTGTGA-GTACCCCCACCCAG 1662	рь
182 182	Qy		141 sLeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSer 156	Qy
2622 TTTTGTATTTTTAGTAGAGACGGGGTTTCACCGTGTTAGCCAAGATGGTCTTGATCACCT 2681	рь			DЬ
182 182	Qy	-	_	Qy
2562 CCTGCCTCAGCCTCCCAAGTAGCTGGGACTACAGGAGCCCACCACCACCACCCCGGCTAATT 2621	Db			Db
	Qy			VΩ
2502 GAGTGCAGTGGCACGATCTTGGCTCACTACAACCTCTGCCTGGGTTCAAGTGATTCT 2561	Db		TCCCA	Дb
182 182	Qγ	-		Qy
2442 CTTTTTTTTTTTTTTTTTTTTTTTTTTGAGATGGAGACTTGCTCTGTCACCCAGGCTG 2501	DЪ		1365 TGTCTTTCGCCTATTTCTGTCTCCATTATCTTGGGATAATGCCTCTGCCTCTCCATGGGA 1424	υр
182 182	Qy		97 97	Qy
2382 TTAGCAGCTAATTCACAGGCCTTTGATCAGAGGTAAGGGACTTTCTGTAGCTATTCAAGT 2441	Db		1305 GCCTCTGGTCGGCCTTAATCTCTGAGCTTCTCTCTTTTTAGGGTAGCCCTGTACCTGTC 1364	р
182 182	Qy		97 97	Qy
2322 TGGGCAAGTGCAGAATTCGCGTCTCTGATGTGTGTGTTCCCTTTCTCAACATGGGATG 2381	Ф		1245 TCAGGGTGCCCTTGCCCCTTCTCTCTCTGGTGACCTTCCCATCTCTCCATGTGCCTTG 1304	В
182 182	Qy		97 97	Оy
2262 AAATAAATAAATACAAAGGGCTGAGAGTCAGGACTGTGCTGCTAGTTCTCTAGGGGATCT 2321	₽		1185 AACTTGGGGTGTGAGGGTCCTGCACCCACCCTTGCCAGAACCCTCCACCCTGATCCTCCT 1244	ф
182 182	Qy		97 97	Qy
2202 AGACCTATTAAATAAATAATAATATTAAATCTATTAAATATTAAATATTAAATCTATT 2261	ДĎ		1125 TGCTTGGCAGGGAGTTCCTAAGGACAGGCCTTTCTGAAGGAAG	Db
182 182	Qy		97 97	Qy
2142 AGGTTGAGGCTGCAGTGAGCTATGATCGTGCCACTGCACTTCAGCCTGGGCAACAGAGCG 2201	Дb		1065 CCATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGGTGAGGGGCTTCTCAG 1124	타
182 182	Qy		84 sHisAsnSerGluCysAlaArgCysGlnAlaCysAspGlu97	Qy
2082 GATGATTTGTCACTTACACAGAGAGTCACATTGATATAGCTTTAAAACTTGGGCTGAAGG 2141	ΔQ		1005 CTGCGGCAACTCCACCTGCCTTGTGTCCCCAAGACACCTTCTTGGCCTGGGAGAACCA 1064	Db
182 182	Qy		64 oCysGlyAsnSerThrCysLeuValCysProGlnAspThrPheLeuAlaTrpGluAsnHi 84	Qy
2022 AGGGTATTTGCCTTCACTTTCAGCCAGCATTCCCCCCCAGCGATCCTAGCCCAGATATTACA 2081	Db		945 GCCTGCCTGACCCTGCCTGGTTCCACAGGGCACTACCTGAAGGCCCCTTGCACGGAGCC 1004	Db ·
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1962 CACTTCAGGCAAACTTAGATGGCAAAAGAGCAATCTGGATCCGCCCTTAGCCAGATACATA 2021	ДЬ		885 GAGAAAGAGGGAGGGAGGGCAGGGTGGGAAGCAGGTCGGGGGTTGCTGGGCAGCCAGC	Db
182 182	Qy		53	Qy
1902 TTCAGGAATAGGAAGAGGGGATAGGGAGGAGGGAGCCTTGGCCCCTGTGATGGGTGGG	뫄		825 AGGTAAGAGGAGGCTGGCAGGCGAGGTAGGGGTAGGCTGACAGAGAAGTAGGGGAGCTGGA 884	Db
182 182	Qy	-		Qy
1842 GATGGAGGGAAGGGCGGCTGGGAGCAGGAGCAGGGGCCTGGGGTGGGGCAGGTGCTGCTGG 1901	Db		765 CCACAGGGGTGGGAGAGGCATGGGGGCAGGCAGGGCTGGAGAGGTGGCGGGCAGGCCCGGG 824	Db
182 182	Qy		53 53	Qy
1783 CTTCTATGAACATGGCGATGGCTGCGTGTCCTGCCCCACGTA-ATTCCTAGCTGTCGTGG 1841	B 1		3/ GIYASPPNEHISLYSIJSIIeGlyLeuPheCysCysArgGlyCysProAla 53	da V
wPheTvrGliHideGlvAsnClvCvsValucovnonnantavionCibCCibCCibCCibCCibCCibCCibCCibCCibCCib	0 <b>v</b>		2	? 5
1723 ATGCCTCTCCTGGATTGCAGGTTCCCGCAGAGATACTGACTG	Db	_	17 LeuValLeuLeuGlyAlaArgAlaGlnGlyGlyThrArgSerProArgCysAspCysAla	Ş Ş
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	3940	AGAGCAGGGCCAACTGTGGGAGGTGTCATTGCGGGGGGTGTCTGACTGA	3881	밁
	247		247	δô
	3880	AGGAAGGAGGCGGGGCAGGGGACTCAAGGCAGAAGCAAGAGTTCTGCTGGGCTACAGTG	3821	В
	247		247	Q
	3820	ATTCGTACACAGGAGCTAACAGTTCAATGGAAGGAGAGCCCCATGGTGCTGGGGGACAAG	3761	Вb
	247		247	οy
. <u>:</u>	3760	CAGTGGATGAGACCAGCACAGTGCCTGCCCTCAAGGGGGTGCTCAGTCAG	3701	В
	247		247	Qy
	3700	CTTGGCACGCGGCCAGCACCGGGTAGCCCTAGTGGACAGAGGTGTTGGGAGCAGAGTCAT	3641	Db
	247		247	Qy
	3640	TGTGTGATTCTGGGCTGCCTTCTGGAGCTGGAAGATCAAGCCTTACTATGATCCCTGGAG	3581	рь
	247		247	Qy
	3580	GTCTGTCAGCAGATGAAGCTGGGATGGAGGCTCTGACCCCCACCACCGGTAAGAACCTCAC	3521	В
	247	laAspGluAlaGlyMetGluAlaLeuThrPr	236	Qy
	3520	GAGCTGTACCAGCACCCAGGTCCAGGAGGCTTGCCTGGTGGCTGACCGCAATCTCTCTGT	3461	Дb
	235		235	Qy
	3460	AACCCTGATACAGAAGGGGAAACTGAGGCAGGGAGTGTGGGGGTGCAGAGGAACCCCTAGAG	3401	В
	235		235	Qy
	3400	CACACACCCACACACGCACCCAGAAGCCTGGGGTCAGGATGGGTAGCCCAGAGTCTACTC	3341	В
	235		235	Qy
	3340	CACCCTGACCTACACATACCGCCACTGCTGGCCTCACAAGCCCCTGGTTACTGGTAAGTA	3281	Ф
	235	aThrLeuThrTyrThrTyrArgHisCysTrpProHisLysProLeuValThr	218	Qy
	3280	TCAGTG -TTCTGGGTCCAGGTGCTCCTGGCTGGCCTTGTGGTCCCCCTCCTGCTTGGGGC	3222	DЬ
	218	-GlnMetPheTrpValGlnValLeuLeuAlaGlyLeuValValProLeuLeuLeuGlyAl	199	Ωy
	3221	GATGGACCGAGGGGGGGGTGAGGAGGGGGGGCAACCACCCAACACCCACC	3162	皮
	198		198	Qy
	3161	CGCTGTCTGTGGCTGGAGGCAGAGTAGGTGGTGGTGCTGGGAATGCGAGTGGGAGAACTGG	3102	Дb
	198	aAlaValCysGlyTrpArg	192	Qy
	3101	CCCTCAGGCCACCCCGTCCCTTGCAGGAGCACCCTGGGGAGCTGTCCAGAGCGCTGTGC	3042	DЬ
	192	ThrLeuGlySerCysProGluArgCysAl	183	Qy
	3041	CTAGGATGGAGCAGATGCTGGGCAGTCTTAGGGAGCCCCTCCTGGCATGCACCCCCTCAT	2982	Db
	182		182	Q
	2981	AGGGACACAATGACAGGAACATTCTTGGGTAGAGTGAGAGGCCTGGGGAAGGGAAGGGTCT	2922	망
	182		182	Qy
	2921	TCTAGTGACTCTTCATGTAGTCTAGAAGAAGCTCAGTGAATAGTGTCTGTGGTTGTTACC	2862	₽
	182		182	γQ
	2861	GTGCTGGGGATGCAACAGGGGACAAAATAGGCAAAATCCCTGTCCTTTTGGGGTTGACAT	2802	В

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4720	61 CAGCAGCCCGCGGGCCTCGGAGCCGTTTACGCGGCCCTGGAGCGCATGGGGCTGGACGGC	46	망
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4660	601 GAGGTGGAGATCGGCCGCTTCCGAGACCAGCAGTACGAGATGCTCAAGCGCTGGCGCCAG	Δ.	뫄
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4600	541 CGGCGCTGGAAGGAGTTCGTGCGCACGCTGGGGCTGCGCGAGGCAGAGATCGAAGCCGTG	-	d d
364			δõ
4540	.481 CCAGCCATGATGCTGCAGCCGGGCCCGCAGCTCTACGACGTGATGGACGCGGTCCCCAGCG	_	ďď
344			γQ
4480	421 GGCCCTGCCCCAGGCCCCGCTGCTGCCGCCACACTCTCGCCAGAGTCCCCAGGCCGGCTCG		망
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4420	361 CACGAACGCCCCTGACTCTGCCTCCCGACCGGGCCCACGTACCCCAATTGGCTCTCTCT	•	당
308	308		ρ
4360	4301 CGGAGTCCGCCCTATGCCCTGACCCACCGGATCCAGCGGGCTTCAGCCCTGGGGTACCCCG	_	ద
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4300	241 TCCTGTCTGCGGTGGGAAGTTGTGGTTTCACAACGTGTTCCCCTTTCTGCCCCCTAACTGA	_	дg
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4180	4121 ACCCCTGGCTACCCCGGAGACCCAGGAGGCGCTCTGCCCCGCAGGTGACATGGTCCTGGGAC	_	밁
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4120	061 CTAGCACCTCCTGACAGCAGTGAGAAGATCTGCACCGTCCAGTTGGTGGGTAACAGCTGG	_	뫄
281			δō
4060	001 GGGCCCCTCTTGGCTTCCAGGCCACCCATCTGTCACCCTTGGACAGCGCCCACACCCTT	-	닭
261	-AlaThrHisLeuSerProLeuAspSerAlaHisThrLeu		γo
4000	941 TGTCCCCTCCTGGAGAGGCACTGCGGGTAAGGGGCCTTACTTGGCAAGCAGGGCTGACCT	w	DЬ
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GenCore version 5.1.4\_p5\_4578 Copyright (c) 1993 - 2003 Compugen Ltd.

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-TRANS-human40.cdi -LIST=45 -DOCALIGN=200 -THR_SCORE-pct -THR_MAX=100
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-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPDF=10 -XGAPEXT=0.5 -FGAPDP=6
-FGAPEXT=7 -YGAPDP=10 -YGAPEXT=0.5 -DELLDP=6 -DELEXT=7
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-MODEL-frame+_p2n.model -DEV-xlh
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/cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/PCTUS_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
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/cgn2_6/ptodata/1/pubpna/PCT_NEW_PUB.seq:*
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/cgn2_6/ptcodata/1/pubpna/US10_NEW_PUB.seq:*
/cgn2_6/ptcodata/1/pubpna/US10_NEW_PUB.seq:*
/cgn2_6/ptcodata/1/pubpna/US60_NEW_PUB.seq:*
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4 9 US-10-081-280-9
4 9 US-10-112-793-9
4 9 US-10-112-193-11
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Sequence 9, Appli
Sequence 9, Appli
Sequence 11, Appl
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Sequence 1, Appli Sequence 14, Appl Sequence 41, Appl	1, 7	Sequence 11, Appl Sequence 11, Appl	11,	N	Sequence 1, Appli	16, 1	26,	16, App	236	֓֞֞֜֜֜֝֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֡֓֜֝֓֓֓֓֡֓֜֝֡֓֡֓֡֡֝֡֓֡֓֡֓֜֝֡֡֡֡	_ ,	Sequence 1, Appli	_ [	14, 7	24, App	14,	_	1601, A	ν,	e 2	, v	Sequence 2, Appli	ູ້	ຸ່ດ	e 5,	5, 1	Ç,	5, Apr	e 1	1577,	1389	۳.	Sequence 9, Appli

## ALIGNMENTS

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US-09-333-966-3
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                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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APPLICANT:
                                                                                                                                                                                                                                                  APPLICANT: Gentz, Reiner L.
APPLICANT: Dillon, Patrick J.
TITLE OF INVENTION: Death Dom
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
                                                          COUNTRY: USA
ZIP: 20005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                 CURRENT APPLICATION DATA:
                                                                                                                                                                                                  STREET: 1100 New CITY: Washington
CLASSIFICATION:
                                 APPLICATION NUMBER:
                                                                                                                                                                                       STATE:
                                                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                               , Application US/09333966
US20020009773A1
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                                                                                                                                                                                                                                                                                                                           Dixit, Vishva
Gentz, Reiner
                                                                                                                                                                                                                                                                                                                                                         Yu, Guo-Liang
Ni, Jian
                                                                                                                                                                                                                                                                           Death Domain : 17
                                 US/09/333,966
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Best Local Similarity:
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APPLICATION NUMBER: US 60/028,711

FILING DATE: 17-OCT-1996

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 60/013,285

FILING DATE: 12-MAR-1996

ATTORNEY/AGENT INFORMATION:

NAME: Steffe, Eric K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 202-371-2540
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 14/
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-771-2600
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       161
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APPLICATION NUMBER:
FILING DATE:
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TYPE: nucleic acid
STRANDEDNESS: double
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                                CysLeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThr
                                                                 AspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysPro
                      TGCCTAGACTGCGGGGCCCTGCACCGCCACACGCCTACTCTGTTCCCGCAGAGATACT
                                                                                                              CAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCCAGGC
                                                                                                                          GlnValAlaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGly
                                                                                                                                                           TGGGAGAACCACCATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGCCTCC
                                                                                                                                                                       TrpGluAsnHisHisAsnSerGluCysAlaArgCysGlnAlaCysAspGluGlnAlaSer
                                                                                                                                                                                                      GGGGCCCGGGCCCAGGCGCACTCGTAGCCCCAGGTGTGACTGTGCCGGTGACTTCCAC
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Matches:
Conservative:
Mismatches:
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RESULT 2
US-10-081-280-9
; Sequence 9, Application US/10081280
; Patent No. US20020165157A1
; GENERAL INFORMATION:
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                                                                               APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-2 LI AND AJ
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
CORRESPET: 460 Point San Bruno B:
CTTY: South San Francisco
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     COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb f
COMPUTER: 1EM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA;
                                                                                                                                                                                                                      GGGCTGGACGGCTGCGTGGAAGACTTGCGCAGCCGCCTGCAGCGCGGCCCG
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                                                                 STATE: California COUNTRY: USA
APPLICATION NUMBER:
                                                         ZIP: 94080
US/10/081,280
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Query
DB:
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Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 1634 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: Linear SEQUENCE DESCRIPTION: SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-5416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CysThrGluProCysGlyAsnSerThrCysLeuValCysProGlnAspThrPheLeuAla
                                                                                                                                                                                                                                                                                                                                                                                                                  LysLysIleGlyLeuPheCysCysArgGlyCysProAlaGlyHisTyrLeuLysAlaPro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GlyAlaArgAlaGlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHis
  PheTrpValGlnValLeuLeuAlaGlyLeuValValProLeuLeuLeuGlyAlaThrLeu
                                         ThrSerThrLeuGlySerCysProGluArgCysAlaAlaValCysGlyTrpArgGlnMet
                                                                                            AspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysPro
                                                                                                                                   TGCCTAGACTGCGGGGCCCTGCACCGCCACACGGCTACTCTGTTCCCGCAGAGATACT
                                                                                                                                              CysLeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThr
                                                                                                                                                                                                                                                                                                                                                                                                                                                       GGGGCCCGGGCCCAGGCACTCGTAGCCCCAGGTGTGACTGTGCCGGTGACTTCCAC
                                                                                GACTGTGGGACCTGCCTGGCTTCTATGAACATGGCGATGGCTGCGTGTCCTGCCCC
                                                                                                                                                                                                                                                                                                                                                 TGCACGGAGCCCTGCGCAACTCCACCTGCCTTGTGTGTCCCCAAGACACCTTCTTGGCC
                                                                                                                                                                                                                                                                                                                                                                                                     AAGAAGATTGGTCTGTTTGCTGCAGAGGCTGCCCAGCGGGGCACTACCTGAAGGCCCCT
                             TELEFAX: 415/952-9881
TELEX: 910/371-7168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007R1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 21 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION DATA:
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                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-2 LI
NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, In
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                                                                                                                      ZIP: 94080

COMPUTER REALABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE WINPATH (Genentech)
                                                                PRIOR APPLICATION DATA:
                                                                                                                    CURRENT APPLICATION DATA:
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                                                                            APPLICATION NUMBER: US/10/112,793 FILING DATE: 28-Mar-2002 CLASSIFICATION: <Unknown>
APPLICATION NUMBER: US/08/828,683A FILING DATE: 31-Mar-1997 APPLICATION NUMBER: 08/625328 FILING DATE: 1-Apr-1996 APPLICATION NUMBER: 08/710802
                                                                                                                                                                                                                          CITY: South San Francisco
STATE: California
                                                                                                                                                                                                               COUNTRY:
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US-09-993-234-6 (1-417) x US-10-112-793-9 (1-1634)
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STRANDEDNESS: Single
TOPOLOGY: Linear
SEQUENCE DESCRIPTION: SEQ ID
US-10-112-793-9
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ATTORNEY/AGENT INFORMATION:
NAME: MAISCHARG, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
ThrTyrThrTyrArgHisCysTrpProHisLysProLeuValThrAlaAspGluAlaGly
                                  ThrSerThrLeuGlySerCysProGluArgCysAlaAlaValCysGlyTrpArgGlnMet
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Publication No. US20030004313A1
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                                    APPLICATION NUMBER: 08/928,069
FILING DATE: 11-Sep-1997
APPLICATION NUMBER: 60/026943
FILING DATE: 23-Sep-1996
ATTORNEY/AGENT INFORMATION:
NAME: Marschang, Diane L.
                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
       NAME: MATSCHANG, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1052R1
TELECOMMUNICATION INFORMATION:
                                                                                                              PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                          APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-3 POLYPEPTIDE NUMBER OF SEQUENCES: 15
                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                               GGGCTGGACGGCTGCAAGACTTGCGCAGCCGCCTGCAGCGCGCCCG 1339
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                                                                                                                                                                                                                                        ADDRESSEE: Genentech, Inc.
STREET: 1 DNA WAY
CITY: South San Francisco
STATE: California
COUNTRY: USA
                                                                                                                      APPLICATION NUMBER: US/10/112,193
FILING DATE: 28-Mar-2002
CLASSIFICATION: <Unknown>
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Percent Similarity:
Best Local Similarity:
Query Match:

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Query Match: DB:

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LENGTH: 1634 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
SEQUENCE DESCRIPTION: SEQ ID
US-10-112-193-11
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Best Local Similarity:
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INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
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LeuLeuAlaProProAspSerSerGluLysIleCysThrValGlnLeuValGlyAsnSer
                                             MetGluAlaLeuThrProProProAlaThrHisLeuSerProLeuAspSerAlaHisThr
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Matches:
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US-09-993-234-9
; Sequence 9, Application US/09993234
; Patent No. US20020146768A1
; GENERAL INFORMATION:
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                                                   TELEFAX: 415/952-9881
TELEX: 910/371-7168
INFORMATION FOR SEG ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 1634 base pairs
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APPLICATION NUMBER: 08/828,683
FILING DATE: CUDANOWN>
ATTORNEY/AGENT INFORMATION:
NAME: MATSCHAMS, Diame L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007P1
                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/993,234
FILING DATE: 19-NO. US2002014676BA1-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Ashkenazi, Avi
TITLE OF INVENTION: Apo-2
NUMBER OF SEQUENCES: 11
SEQUENCE
                                                                                                                            TELECOMMUNICATION INFORMATION: TELEPHONE: 415/225-5416
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CITY: South San Francisco
STATE: California
                         TYPE: Nucleic Acid
STRANDEDNESS: Single
               TOPOLOGY: Linear
DESCRIPTION: SEQ
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                                                                                                    Asp {\tt GlnLeuProSerArgAlaLeuGlyProAlaAlaAlaProThrLeuSerProGluSer}
              TGGACCCCTGGCTACCCCGAGACCCAGGAGGCGCTCTGCCCGCAGGTGACATGGTCCTGG
                                          CTTCTAGCACCTCCTGACAGCAGTGAGAAGATCTGCACCGTCCAGTTGGTGGGTAACAGC
                                                  LeuLeuAlaProProAspSerSerGluLysIleCysThrValGlnLeuValGlyAsnSer 280
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Matches:
Conservative:
Mismatches:
Indels:
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FILING DATE:

CLASSIFICATION:

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/08/815,469

FILING DATE:

APPLICATION NUMBER: NO. US20020009773A1 Yet Assigned

APPLICATION DATA:

APPLICATION DATA:

APPLICATION NUMBER: US 60/028,711

FILING DATE: 17-OCT-1996

PRIOR APPLICATION DATA:

ADDITION TOWNSER: US 60/028,711

FILING DATE: US/0728,711

FILING DATE: US/0728,711
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Patent No. US20020009773A1
GENERAL INFORMATION:
APPLICANT: Yu, Guo-Liang
                                     TELEFAX: 202-371-2540 INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 1783 base pair
TYPE: nucleic acid
                                                           NAME: Steffe, Eric K.
REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 14
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
                                                                                                                                                                                                                                                                                                                                 ZIP: 20005-3934

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
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APPLICANT:
APPLICANT:
                                                                                                                               APPLICATION NUMBER: US 60/013, FILING DATE: 12-MAR-1996 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Gentz, Reiner L.
APPLICANT: Dillon, Patrick J.
TITLE OF INVENTION: Death Domain
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
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STREET: 1100 New
CITY: Washington
STATE: DC
COUNTRY: USA
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1100 New York Ave.,
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Ni, Jian
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NW, Suite
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TOPOLOGY: both
MOLECULE TYPE: C
FEATURE:
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                              GATGAAGCTGGGATGGAGGCTCTGACCCCACCACCGGCCACCCATCTGTCACCCTTGGAC
                                     AspGluAlaGlyMetGluAlaLeuThrProProProAlaThrHisLeuSerProLeuAsp
                                                            GGGGCCACCCTGACCTACACATACCGCCACTGCTGGCCTCACAAGCCCCTGGTTACTGCA
                                                                                          TGGAGGCAGATGTTCTGGGTCCAGGTGCTCCTGGCTTGGCCTTGTGGTTCCCCCCTCCTGCTT
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PRIOR APPLICATION NUMBER: US/60/234,034
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PRIOR APPLICATION NUMBER: US/60/234,567
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NUMBER OF SEQ ID NOS: 1392
SOFTWARE: PatentIn version
SEQ ID NO 1389
LENGTH: 1743
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                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1389, Application Patent No. US20020165180A1
                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Process for Identifying TITLE OF INVENTION: Gene Sets FILE REFERENCE: 689290-77 CURRENT APPLICATION NUMBER: US/09/954,531 CURRENT FILING DATE: 2002-05-02
                                                                                                                                         TYPE: DNA
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MetMetLeuGlnProGlyProGlnLeuTyrAspValMetAspAlaValProAlaArgArg
                GAGACCCAGGAGGCGCTCTGCCCGCAGGTGACATGGTCCTGGGACCAGTTGCCCCAGCAGA
                                             GluThrGlnGluAlaLeuCysProGlnValThrTrpSerTrpAspGlnLeuProSerArg
                                                            CysTrpProHisLysProLeuValThrAlaAspGluAlaGlyMetGluAlaLeuThrPro 246
                                                                                                                              TGCTGGCCTCACAAGCCCCTGGTTACTGCAGATGAAGCTGGGGATGGAGGCTCTGACCCCA
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Oy 347  Db 1208  Oy 367  Db 1268  Qy 387  Db 1328  Qy 407  Db 1328  RESULT 8  US-10-092- Sequence FILE REI GENERAL HAPPLICAL HITLE REI CURRENT						,,					•			•	•										
TrpLysGluPheValArgThrieuGlyLeuArgGluAlaGluIleGluAlaValGluVal 3	δ	Ъ	Qy	Db	Qy	3	, Oy	dd Qy	Дy	9 dq		Ali Pre Sco Per Bes Que DB:					RES US: ; S	DЬ	Qy	סט גי	, Db	Qy	망	Qy	Db
76 - 23 - 34 - 34 - 34 - 34 - 34 - 34 - 34	54GlyHisTy 	5484 GAGAAAGAGGGAGGGAGGGCAGGGTGGAAAGCAGGTCGG	53	5424 AGGT	Qy 53	, Db 5364 CCACAGGGGTGGG	(Oy) 53	Db 5304 G	Qy 17 Db 5244	3 GlnärgProärgGlyCys :::           5184 GAGGCCCCACGTGGGCAGCTGCGCTGAGAGTCAGCA	-09-993-234-6 (1-417) x US-10-092-154-1577 (1-10797	Scores:       4.34e-129       Length:       1079	SEQ ID NO 1577 LENGTH: 10797 TYPE: DNA ORGANISM: Homo sapiens -10-092-154-1577	Prior Application removed - See File Wrapper or Pa SOFTWARE: PatentIn Ver. 2.0	CURRENT APPLICATION NUMBER: US/10/092, CURRENT FILING DATE: 2002-03-07	APPLICANT: Rosen et al. TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodi EILE REFERENCE: PC009C1	SULT 8 -10-092-154-1577 -10-092-154-1577 Sequence 1577, Application US/100921 Publication No. US20030054375A1 GENERAL INFORMATION:	1388 GAAGACTTGCGCAGCCGCCTGCAGCGTGGCCCG 142	407	1328	1268	367	1208	347	Db 1148 ATGATGCTGCAGCCGGGCCCGCAGCTCTACGACGTGATGGACGCGGTCCCAGCGCGGCGC
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Qy 192 aAlaValCysGlyTrpArg198	. :	182
		6561 CACTTCAGGCAAACTTAGATGGCAAAAGAGCAATCTGGATCCGCCTTAGCCAGATACATA 6620
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Qy 182 182		
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Qy 182 182		
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Qy 182 182		vPheTvrGluH1sGlvAspGlvCvsValSerCvsProThrSer
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Qy 182 182		ArgArgAspThrAspCvsGlvThrCvsLeuProGl
Db 7341 GCGCCCGGCCTCCATTCAAGTCTTTATTGAATATCTGCTATGTTCTACACACTGTTCTAG 7400		GGCTCTCTACTCCCAGACCCCCTTCTCCCTGCCTGACCCACTCCTGTCCCATGGTGACGC
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Db 7281 GACCTCGTGATCCACCCGCCTTGGCCTCCCAAAGTGCTGGGATTACAGGCATGAGCCACC 7340		
Qy 182 182		sLenAspCvsG1vA1aLenHisArgHisThrArgLenLenCvsSer
Db 7221 TTTTGTATTTTTAGTAGAGACGGGGTTTCACCGTGTTAGCCAAGATGGTCTTGATCACCT 7280		
Qy 182 182		pPheValGluCysGlnValSerGlnCysValSerSerSerProPheTvrCvsGlnProCv
Db 7161 CCTGCCTCAGCCTCCCAAGTAGCTGGGACTACAGGAGCCCACCACCACCCCCGGCTAATT 7220		
Qy 182 182		101 nValAlaLeuGluAsnCvsSerAlaValAlaAsnThrArgCvsGlvCvsLvsDroGlvTr
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Qy 182 182		97 97
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Qy 182 182	·	97
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Qy 182 182	<del></del>	97
Db 6861 AAATAAATAAATACAAAGGGCTGAGAGTCAGGACTGTGCTGCTAGTTCTCTAGGGGATCT 6920	·	AACTTGGGGTGTGAGGGTCCTGCACCCACCCTTGCCAGAACCCTCCACCCTGATCCTCCT
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Db 6801 AGACCTATTAAATAAATAAATAATATTAAATCTATTAAATATTAAATATTAAATCTATT 6860		TGCTTGGCAGGGAGTTCCTAAGGACAGGCAGGCAAAGTTGGCAAGCTGGCTTGGCTCGCGCCCA
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	AGCAGGGCCAACTGTGGGAGGTGTCATTGCGGGGGTGTCTGCTGACTGA	8480 AG	DЪ
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	GTACACAGGAGCTAACAGTTCAATGG	8360 ATTC	Db
	247	247	Qy
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	CCGGTAAGAACCTCAC 8	8120 GT	Db
	AspGluAlaGlyMetGluAlaLeuThrProF	236	Qy
	CTGTACCAGCACCCAGGTCCAGGAGGCTTGCCTGGTGGCTGACCGCAATCTCTCTGT 8	8060 GAG	B
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	CCTGATACAGAAGGGGAAACTGAGGCAGGGAGTGTGGGGTGCAGAGGAACCCTAGAG 80	8000 AAC	Дb
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	ACACACCCACACACGCACCCAGAAGCCTGGGGTCAGGATGGGTAGCCCAGAGTCTACTC 7999	7940 C	Ъ
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	ACCCTGACCTACACATACCGCCACTGCTGGCCTCACAAGCCCCTGGTTACTGGTAAGTA 7939	7880 c	DЬ
	LeuThrTyrThrTyrArgHisCysTrpProHisLysProLeuValThr 23	218 a	Qy
	<b>8</b> 0 H	7821 TC	DЬ
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3 GlnArgProArgGly	Alignment Scores:  Pred. No.:  10797  Score:  Percent Similarity:  Percent Similarity:  109.038  Query Match:  1008  100	RESULT 9 US-09-764-847-1577 Sequence 1577, Application US/09764847 Patent No. US20020132767A1 GENERAL INFORMATION: APPLICANT: Rosen et al. FILE OF INVENTION: Nucleic Acids, Proteins, and Antibodi FILE REFERENCE: PC009 CURRENT APPLICATION NUMBER: US/09/764,847 CURRENT FILING DATE: 2001-01-17 Prior application data removed - consult PALM or file wra NUMBER OF SEQ ID NOS: 2003 SOFTWARE: Patentin Ver. 2.0 SEQ ID NO 1577 LENGTH: 10797 TYPE: DNA ORGANISM: Homo sapiens	Db 926 Qy 40 Db 932	Qy 345  Db 9140  Qy 365  Db 9200  Qy 385	309	308	308	Db 8840 TCCTGTCTGCGGTGGGAAGTTGTGGTTTCACAACGTGTTCCCCTTTCTGCCCCCTAACTGA
		pper	euAspGly          TGGACGGC	3luAlaVal	AlaGlySer           GCGGCTCG     AlProAla 	GCTCTCTCT	GGGTACCCG	CCTAACTGA
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PheValGluCysGlnValSerGlnCysValSerSerProPheTyrCysGlnProCy 
ValAlaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysp 
CCTTTGGCCCTGACTAACTCTCCCACTCCCCATCTCCCTGCACCCCCACCAGCC
CCA
CCTCTGGTCGGCCTTAATCTCTGAGCTTCTCTCTTTTTTAGGGTAGCCCTGTACC
AGGGTGCCCTTGCCCTTCTCTTTCC
TGGGGTGTGAGGGTCCTGCAC
TGCTTGGCAGGGAGTTCCTAAGGACAGGCCTTTCTGAAGGAAG
HisasnSergluCysalaargCysGlnalaCysaspGlu- 
oCysGlyAsnSerThrCysLeuValGysProGlnAspThrPheLeuAlaTrpGluAsn 
CCTGGTT
AGAAAGAGGGAGGGAGGGCAGGGTGGAAAGCAGGTCGGGGGTTGCTGGGCAGCCCC
AGGTAAGAGGAGGCTGGCAGGGGAGGTAGGGGTAGGCTGAC
CCACAGGGGTGGGAGAGGCATGGGGCAGGC

7221 TTTTGTATTTTAGTAGAGACGGGGTTTCACCGTGTTAGCCAAGATGGTCTTGATCACCT 182 7281 GACCTCGTGATCCACCCGGCCTTGGGCCTCCCAAAGTGCTGGGATTACAGGCATGAGCCACC 182	0 0 0 0	
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182	Qy	
6741 AGGTTGAGGCTGCAGTGAGCTATGATCGTGCCACTGCACTTCAGCCTGGGCAACAGAGCG	DЬ	
182	Qy	
6681 GATGATTTGTCACTTACACAGAGAGTCACATTGATATAGCTTTAAAACTTGGGCTGAAGG	문	
182	Qy	
6621 AGGGTATTTGCCTTCACTTTCAGCCAGCATTCCCCCCAGCGATCCTAGCCAGATATTACA	р	
182	Qy	
6561 CACTTCAGGCAAACTTAGATGGCAAAAGAGCAATCTGGATCCGCCTTAGCCAGATACATA	Db	
182	Qy	
6501 TICAGGAATAGGAAGAGGGGATAGGGAGGGAGGCCTTGGCCCCTGTGATGGGTGGG	В	
182	Qγ	
6441 GATGGAGGGAAGGGCGGCTGGGAGCAGAGCAGGGGCCTGGGGTGGGGCCAGGTGCTGCTGG	皮	
182	Qγ	
6382 CTTCTATGAACATGGCGATGGCTGCGTGTCCTGCCCCACGTA-ATTCCTAGCTGTCGTGG	рь	
TCCTGCCCCACG		6382 182 6441

Qy	182		82	-
Ъ	7521	AGGGACACAATGACAGGAACATTCTTGGGTAGAGTGAGAGGCCTGGGGAGGGA	580	
Qy	182	18	32	
Ъ	7581	CTAGGATGGAGCAGATGCTGGGCAGTCTTAGGGAGCCCCTCCTGGCATGCACCCCCTCAT 76	540	:
Qy	183	LeuGlySerCysProGlu	32	
ДЪ	7641		700	
Qy	192	AlaValCysGlyTr	86	
DЬ	7701	CGCTGTCTGTGGCTGGAGGCAGAGTAGGTGGTGCTGGGGAATGCGAGTGGGAGAACTGG 77	60	
Qy	198	19	æ	
Дb	7761	GATGGACCGAGGGGAGGCGGGTGAGGAGGGGGGGGCAACCACCCAACACCCACC	120	
Qу	199	PheTrpValGlnValLeuLeuAlaGlyLeuValValProLeuLeuLeuGlyAl 21	œ	
В	7821	TCAGTG-TTCTGGGTCCAGGTGCTCCTGGCCTTGTGGTCCCCCTCCTGCTTGGGGC 78	179	
Qy	218	ThrLeuThrTyrThrTyrArgHisCysTrpProHisLysProLeuValThr 23	Ö	
В	7880	CTGACCTACACATACCGCCACTGCTGGCCTCACAAGCCCCTGGTTACTGGTAAGTA 79	39	
Qγ	235	23	Ġ	
망	7940	CACACACCCACACACCCAGAAGCCTGGGGTCAGGATGGGTAGCCCAGAGTCTACTC 79	99	
γO	235	23	U)	
Ъ	8000	AACCCTGATACAGAAGGGGAAACTGAGGCAGGGAGTGTGGGGTGCAGAGGAACCCTAGAG 80	.59	
Qγ	235	23	Сī	
Ъ	8060	GAGCTGTACCAGCACCCAGGTCCAGGAGGCTTGCCCTGGTGGCTGACCGCAATCTCTCTGT 811	19	
Qy	236	spGluAlaGlyMetGluAlaLeuThrProPro 24	7	
Db	8120	CAGATGAAGCTGGGATGGAGGCTCTGACCCCACC	79	
VΩ	247	24	7	
ఠ	8180	TGTGTGATTCTGGGCTGCCTTCTGGAGCTGGAAGATCAAGCCTTACTATGATCCCTGGAG 823	39	
Qy	247	24	7	
Db	8240	CTTGGCACGCGGCCAGCACCGGGTAGCCCTAGTGGACAGAGGTGTTGGGAGCAGAGTCAT 82	99	
Qy	247		7	
Ъ	8300	CAGTGGATGAGACCAGCACAGTGCCTGCCCTCAAGGGGTGCTCAGTCAG	59	
Qγ	247	24:	7	
В	8360	ATTCGTACACAGGAGCTAACAGTTCAATGGAAGGAGAGCCCCATGGTGCTGGGGGACAAG 841	19	
Ϋ́	247	247	7	
용	8420	AGGAAGGAGGCGGGGCAGGGGACTCAAGGCAGAAGCAAGAGTTCTGCTGGGCTACAGTG 847	79	
Ωy	247	247	7	
용	8480	AGAGCAGGGCCAACTGTGGGAGGTGTCATTGCGGGGGTGTCTGCTGACTGA	39	
γ	247	247	7	
융	8540	TGTCCCCTCCTGGAGAGGCACTGCGGGTAAGGGGCCTTACTTGGCAAGCAGGGCTGACCT 859	99	

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, and a second s	Ωу	D 49	р 2	₽ 2	ДЪ	B 7	용	Qy	Db	QΥ	В	Qy	Db	Qy	P 64	, p	B	Qy
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T 10 -081-280-5 -081-280-5 -081-280-5 ent No. US20020165157A1 ENERAL INFORMATION: Apo-2 LI AND Apo-3 POLYPEPTIDES TITLE OF INVENTION: Apo-2 LI AND Apo-3 POLYPEPTIDES NUMBER OF SEQUENCES: 11 CORRESPONDENCE ADDRESS: ARDDRESSE: Generatech, Inc. STREET: 460 Point San Bruno Blvd CITY: South San Francisco STARTE: California COUNTRY: USA ZIP: 94080 COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Generatech) CURRENT APPLICATION DATA:	CysValGluAspLeuArgSerArgLeuGlnArgGlyPro 417 	GINGLNProAlaGlyLeuclyAlaValTyrAlaAlaLeuGluArgMetGlyLeuAspGly 404 	GluValGluIleGlyArgPheArgAspGlnGlnTyrGluMetLeuLysArgTrpArgGln 384	ArgArgTrpLysGluPheValArgThrLeuGlyLeuArgGluAlaGluIleGluAlaVal 364 	ProAlaMetMetLeuGlnProGlyProGlnLeuTyrAspValMetAspAlaValProAla 344 	GGCCCTGCCCAGGCCCCGCTGCTGCCCCACACTCTCGCCAGAGTCCCCAGCCGGCTCG 9079	CGAACGCCCTGACTCTGCCTCCCGACCGCGGGCCCACGTACCCCAATTGGCTCTCTCT	308	CGGAGTCCGCCCTATGCCCTGACCCACCGGATCCAGCGGGCTTCAGCCCTGGGGTACCCG 8959	308	TCCTGTCTGCGGTGGGAAGTTGTGGTTTCACAACGTGTTCCCCTTTCTGCCCCCCTAACTGA 8899	308	TTC 8	lnLeuProSerArgAlaLeu 308	ThrProGlyTyrProGluThrGlnGluAlaLeuCysProGlnVAlThrTrpSerTrpAsp 301 	LeuA.19ProProAspSerSerGluLysIleCysThrValGlnLeuValGlyAsnSerTrp 281	GGGC	1

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; SEQUENCE DESCRIPTION: SEQ US-10-081-280-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Percent Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Alignment Scores:
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TELEPHONE: 415/225-5416
TELEFAX: 415/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: MAISCHANG, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007R1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                 GlyAlaArgAlaGlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHis
                                                                     AspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysPro 180
                                                                                                                          CysLeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThr
                                                                                                                                                                                   TrpPheValGluCysGlnValSerGlnCysValSerSerSerProPheTyrCysGlnPro
                                                                                                                                                                                                                                         GlnValAlaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGly 120
                                                                                                                                                                                                                                                                                                                                                        CysThrGluProCysGlyAsnSerThrCysLeuValCysProGlnAspThrPheLeuAla
                                                                                                                                                                                                                                                                                                                                                                                                               LysLysIleGlyLeuPheCysCysArgGlyCysProAlaGlyHisTyrLeuLysAlaPro
                       ThrSerThrLeuGlySerCysProGluArgCysAlaAlaValCysGlyTrpArgGln 199
                                                                                                                                                                    TGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGCCAACCA
                                                                                                                                                                                                                            CAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCCAGGC
                                                                                                                                                                                                                                                                                                                                                                                                  AAGAAGATTGGTCTGTTTTGTTGCAGAGGCTGCCCAGCGGGGCACTACCTGAAGGCCCCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                       GGGGCCCGGGCCCAGGGCGCACTCGTAGCCCCAGGTGTGACTGTGCCGGTGACTTCCAC
                                                      GACTGTGGGACCTGCCTGGCTTCTATGAACATGGCGATGGCTGCGTGTCCTGCCCC
                                                                                                             TGCCTAGACTGCGGGGCCCTGCACCGCCACACGGCTACTCTGTTCCCGCAGAGATACT
                                                                                                                                                                                                                                                                                                                                          TGCACGGAGCCCTGCGCAACTCCACCTGCCTTGTGTGTCCCCAAGACACCTTCTTGGCC
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STRANDEDNESS: Single
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/829,270 FILING DATE: 31-Mar-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/10/081,280 FILING DATE: 21-Feb-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 1438 base pairs
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Matches:
Conservative:
Mismatches:
Indels:
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US-10-112-793-5
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Publication No. US20020192729A1
GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-2 LI AND
NUMBER OF SEQUENCES: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                               8
                                                                                                                                                                           557
                                                         497
                                                                                                                     437
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APPLICATION NUMBER: 08/625328 FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-6ep-1996
ATTORNEY/ACENT INFORMATION:
NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007P1
TELEPHONE: 650/225-5416
TELEPHONE: 650/225-5416
                             61
                                                                                     41
                                                                                                                COMPUTER: IBM PC compatible operating SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech) CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: <Unknown> PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 1
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ADDRESSEE: Genentech,
                                                      CysThrGluProCysGlyAsnSerThrCysLeuValCysProGlnAspThrPheLeuAla
TGCACGGAGCCCTGCGGCAACTCCACCTGCCTTGTGTGTCCCCCAAGACACCTTCTTGGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: South San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 1438 base
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Matches:
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Alignment Scores:
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                         Percent Similarity:
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                                                                                                                                                                                                                                                                                          TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 5:
                                                                     No.:
                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 08/928,069
FILING DATE: 11-Sep-1997
APPLICATION NUMBER: 60/026943
FILING DATE: 23-Sep-1996
ATTORNEY/AGENT INFORMATION:
NAME: MATSCHANG, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: p1052R1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                857
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APPLICATION NUMBER: US/10/112/193
FILING DATE: 28-Mar-2002
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                            SEQUENCE
                                                                                                                                                                                                                                                                         SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Apo-3 POLYPEPTIDE NUMBER OF SEQUENCES: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Ashkenazi, Avi
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                                                                                                                                                     TOPOLOGY: Linear NCE DESCRIPTION: SEQ
                                                                                                                                                                                                 TYPE: Nucleic Acid
STRANDEDNESS: Single
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LENGTH: 1438 base pairs
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STREET: 1 DNA Way
CITY: South San Francisco
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                    Length:
Matches:
Conservative:
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US-09-884-733-5
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Query Match:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 5, Application US/09884733
Patent No. US20020123116A1
GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
                                                     ZIP: 94080

COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy di.
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION NUMBER: US/09/884,733
FILING DATE: 19-Jun-2001
CLASSIFICATION: <UNIDADA:
CLASSIFICATION: 
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/304,003
FILING DATE: 14-JUNE-2000
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ADDRESSEE: Genentee
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CITY: South San Francisco
STATE: California
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RESULT 14
US-09-993-234-5
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Best Local Similarity:
Query Match:
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             Sequence 5, Application US/09993234
Patent No. US20020146768A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
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REFERENCE/DOCKET NUMBER: P1007
TELECOMMUNICATION INFORMATION:
  APPLICANT: Ashkenazi, Avi J
                                                                                                                                                                                                                                                                                                                                    GlnValAlaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGly
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TYPE: Nucleic Acid
STRANDEDNESS: Single
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TELEFAX: 415/952-9881
TELEX: 910/371-7168
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FILING DATE: <UOKnown>
ATTORNEY_AGENT INFORMATION:
NAME: MATSCHANG, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE_DOCKET NUMBER: P1007P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-5416
TELEPAX: 415/952-9881
TELEFAX: 415/952-9881
TELEFAX: 910/371-7168
INFORMATION FOR SEQ ID NO: 5:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
                                                                                                                                                                     CysThrGluProCysGlyAsnSerThrCysLeuValCysProGlnAspThrPheLeuAla
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
                                                    GlnValAlaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGly
                                                                                                    TrpPheValGluCysGlnValSerGlnCysValSerSerSerProPheTyrCysGlnPro
                                                                                                                      TrpGluAsnHisHisAsnSerGluCysAlaArgCysGlnAlaCysAspGluGlnAlaSer
                                                                                                                                                                                                                                             AAGAAGATTGGTCTGTTTTGTTGCAGAGGCTGCCCAGCGGGGCACTACCTGAAGGCCCCT
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TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/09/993,234
FILING DATE: 19-No. US20020146768A1-2001
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Conservative:
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APPLICANT: HOTRIGAN, Stephen
TITLE OF INVENTION: Cancer Gene Determination and
TITLE OF INVENTION: Sets
FILE REFERENCE: 689290-73
CURRENT APPLICATION NUMBER: US/09/964,824A
CURRENT FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: US/60/236,033
PRIOR APPLICATION NUMBER: US/60/236,033
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,028
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Best Local Similarity:
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-964-824A-292
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US-09-964-824A-292
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SEQ ID NO 292
LENGTH: 816
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GenCore version 5.1.4\_p5\_4578 Copyright (c) 1993 - 2003 Compugen Ltd.

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Title:
Perfect score:
Sequence:
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-MODEL-frame+_p2n.model -DEV-xlh
-O-/cgn2_1/USPTO_Spool/US09993234/runat_27032003_115456_15362/app_query.fasta_1.2346
-O-/cgn2_1/USPTO_Spool/US09993234/runat_27032003_115456_15362/app_query.fasta_1.2346
-DB-EST -OFMT-fastap -SUFFIX=rst -MINMATCH=0.1 -LOOPCL=0 -LOOPEXT=0
-UNITS-bits -START=1 -END=-1 -MATRIX-blosum62 -TRANS-human40.cdi -LIST=45
-DOCALICN=200 -THR_SCORE-pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15 -MODE-LOCAL
-OUTFMT-pto -NORM-ext -HEAPSIZE=500 -MINLEN=0 -MAXIEN=2000000000
-USER=US09993234_eCGN_11_4749_erunat_27032003_115456_15362 -NCPU=6 -ICPU=3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS=1 -XGAPOF=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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N71141 za80g11.s1	AW182875 xp99a04.x		B212	2d07.r1	42 60278	9 tq18c01.	AV664983 AV664983	p73c0	zd66a06	BI655045 603282719	7014h09	op33h(		BM853061 K-EST0134	٧.	71143 za80h09	Η, (	BE668836 150213 WA	0 N E31003	BM824350 K-FST0095	040 040 N-ESIUU96	BE0900/2 PM4-CT040		W002222	E220347 hv	K-EST001	BM770798 K-EST0054	by08d05.	1e75f02.x	× :	BM756372 K-EST0014	<b>,</b> ,	AW517358 xp93g02.x		w	вм009354 603629813	5 tg19b08.x	0 xv41b12.	66 60133486	1909448 60306	807913 wf52c03.	BQ068309 AGENCOURT	cription

## ALIGNMENTS

AUTHORS TITLE	REFERENCE	ORGANISM	KEYWORDS	VERSION	ACCESSION		DEFINITION	Locus	BQ068309	RESULT 1
NIH-MGC http://mgc.nci.nih.gov/. National Institutes of Health, Mammalian Gene Collection (MGC)	Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi; Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 1010)	Homo sapiens	EST.	BQ068309.1 GI:19897355	BQ068309	5', mRNA sequence.	mo sapiens cDNA clo	BQ068309 1010 bp mRNA linear EST 02-APR-2002		

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              ValCysGlyTrpArgGlnMetPheTrpValGln-ValLeuLeuAlaGlyLeuValValPr
     GTCTGTGGCTGGAGG---
                                                    ASPG1yCysValSerCysProThrSerThrLeuG1ySerCysProGluArgCysAlaAla
                                                                                                          CTCTGTTCCCGCAGAGATACTGACTGTGGGACCTGCCTGGCCTGGCTTCTATGAACATGGC
                                                                                                                                                                                                                    ArgCysGlYCysLysProGlyTrpPheValGluCysGlnValSerGlnCysValSerSer 133
                                                                                                                           LeuCysSerArgArgAspThrAspCysGlyThrCysLeuProGlyPheTyrGluHisGly 173
                                                                                                                                                                                                                                                                                                                                  TCACCCTTCTACTGCCAACCATGCCTAGACTGCGGGGCCCTGCACCGCCACACACGGCTA
                                                                                                                                                                                 SerProPheTyrCysGlnProCysLeuAspCysGlyAlaLeuHisArgHisThrArgLeu 153
                                                                                                                                                                                                                                                                                          AlaCysAspGluGinAlaSerGlnValAlaLeuGluAsnCysSerAlaValAlaAspThr 113
                                                                                                                                                                                                                                                                                                                                                                                          GGCACTACCTGAAGGCCCCTTGCACGGAGCCCTGCGGCAACTCCACCTGCCTTGTGTGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           High quality sequence stop: 689.
Location/Qualifiers
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Contact: Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cloning). Average LIBELL CO. 7-3.5 kb. Library is normal full-length clones and was co (Invitrogen). Research Geneti this is a NIH MGC Library." 9 a 353 c 287 g 191 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5770562"
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Unpublished (1997)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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1 (bases 1 to 698)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 quality sequence stop: 459.
/Clone_lib="Soares_NFL_T_GBC_S1"
//lab_bost="DH10B"
//lab_bost="DH10B"
//note="Organ: pooled; Vector: pT773D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Equal amounts of plasmid DNA from three normalized libraries (fetal lung NbHL19W, testis NHT, and B-cell NCL_CGAP_GCBl) were maxed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was pCR-amplified cDNAs from pools of 5,000 clones made from the same 3 libraries. The pools consisted of 1.M.A.G.E. clones 297480-302087, 68262-687239, 726408-728711, and 729096-731399. Subtraction by Bento Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                                                                         /db_xref="taxon:9606"
/clone="IMAGE:2359204"
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                                              CTGGTGCTGCTGGGGGCCCGGTGCCCAGGGGGCGCACTCGTAGCCCCAGGTGTGACTGTGC
        GCTGTAAGCCAGGCTGGTTTGTGGAGTGCCAGGTCAGCCAGTGTGTCAGCAGTTCACCCT
                   lyCysLysProGlyTrpPheValGluCysGlnValSerGlnCysValSerSerSerProP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies,
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Ma
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     http://image.llnl.gov
Plate: LLAM11538 row: o column:
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5214568"
/clone_lb="NIH_MGC_118"
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/lab_host="DH108"
                                                                                                                                                                                                                                                                                                                                                                                                                 /note="Vector: PCMV-SPORT6; Site_1: NotI; Site_2: ECORV (destroyed); RNA source leukocytes from anonymous pool of non-activated adult donors. Library is oligo-dT primed and directionally cloned (ECORV site is destroyed upon cloning). Average insert size 1.7 kb, insert size range 1.2-3.3 kb. Library is normalized and enriched for full-length clones and was constructed by C. Gruber (Invitrogen). Research Genetics tracking code 027. Note: this is a NIH MGC Library."
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1201.50
92.31%
91.90%
51.72%
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Health, Mammalian
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Matches:
Conservative:
Mismatches:
Indels:
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383 116 263 76 56

143

203

Mismatches: Indels: Conservative:

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194

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BASE COUNT
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High quality sequence stop: 681.
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BE563566.1 GI:
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
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                                                                                                                                                                                                                                                                                                                                                                                                              Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
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601334867F1
                                                                        /clone_ib="NIH_MGC_3y"
/clone_lib="NIH_MGC_3y"
/tissue_type="adenocarcinoma"
/tissue_type="adenocarcinoma"
/lab_host="DH10B (phage-resistant)"
/note="Organ: pancreas; Vector: pOTB7; Site_1: XhoI;
Site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin by Ling Hong in the laboratory of Gerald M. Rubin '"miversity of California, Berkeley) using ZAP-cDNA '"miversity of California, and Superscript II RT (Life
                                                                           110
                                                                       synthes...
Technologies)."
220
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/db_xref="taxon:9606"
/clone="IMAGE:3688721"
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                                                                                                                                                                                                                                                                                                          CAGTACGAGATGCTCAAGCGCTGGCGCCAGCAGCCCGG-GGCCTCGGAGCCGTTTAC
                                                                                                                                                                                                                                                                                                                                                            GlyLeuArgGluAlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGln
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                                               Eukaryota; Metazoa; (Manumalia; Eutheria; Manumalia; Eutheria; 1 to 632)
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                                                                                                                                                                      AW268610 632 bp mRNA linear ESY xv41D12.x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:2815679 3' similar to SW:WSL1_HUMAN Q93038 WSL-1
                   1 (bases 1 to 632)
NCI-GAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy
                                                                                                                           EST
Tumor Gene Index
Unpublished (1997)
                                                                                             Homo sapiens
                                                                                                           human
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Primates;
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Catarrhini; Craniata;

Vertebrata; Euteleostomi; i; Hominidae; Homo.

EST 03-JAN-2000

542

354 482 334

422 314 362 294 302 274 242

602 374

Project (CGAP),

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US-09-993-234-6 (1-417) x AW268610 (1-632)
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   ValSerCysProThrSerThrLeuGlySerCysProGluArgCysAlaAlaValCysGly 196
                                                                                                                                                                                                           CysLysProGlyTrpPheValGluCysGlnValSerGlnCysValSerSerSerProPhe 136
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                                                                                                                                                                                                                                                                                                                                                          ThrPheLeuAlaTrpGluAsnHisHisAsnSerGluCysAlaArgCysGlnAlaCysAsp 96
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CTGGTGCTGCGGGGCCCGGGCCCAGGGCGCACTCGTAGCCCCAGGTGTGACTGTGCC
                                                            ArgArgAspThrAspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCys
                                                                                                                TACTGCCAACCATGCCTAGACTGCGGGGCCCTGCACCGCCACACGCCTACTCTGTTCC
                                                                                                                                   TyrCysGlnProCysLeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSer 156
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                                        CGCAGAGATACTGACTGTGGGACCTGCCTGCCTGGCTTCTATGAACATGGCGATGGCTGT
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Email: cg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This clone is available royalty-free through LINL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Possible reversed clone: similarity on wrong strand possible reversed clone: polyT not found
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note-*Organ: pooled; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Equal amounts of plasmid DNA from three normalized libraries (fetal lung NbHL19W, testis NHT, and B-cell NCI_CGAP_GCBl) were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 297480-302087, 682632-687239, 726408-728711, and 729096-731399. Subtraction by E Soares and M. Fatima Bonaldo.

a 194 c 192 g 143 t 1 others
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/clone="IMAGE:2815679"
/clone=1b="Soares_NFL_T_GBC_S1"
/lab_host="DH10B"
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Insert Length: 2028 Std Error: 0.00
Seq primer: -40UP from Glbco
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Tissue Procurement: Ash Alizadeh,
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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/tissue_type="B-cell, chronic lymphotic leukemia"
/lab_host="DH10B"
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
CDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLN
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LICM1921 row: i column: 04
                                                                                                                                                                 NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
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                                                         e: LLCM1921 row: i column: 04 quality sequence stop: 649. Location/Qualifiers
/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5443659"
/clone_lib="NIH_MGC_41"
/tissue_type="amelanotic m
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107 ø /lab\_host="DH10B (phage-resistant)"
/note="Organ: skin; Vector: pOTB7; Site\_1: xhoI; Site\_2:
ECORI; cDNA made by Oligo-dT priming. Directionally clon
into EcoRI/XhoI sites using the following 5' adaptor:
GGCACGA6(G). Library constructed by Ling Hong in the
laboratory of Gerald M. Rubin (University of Callifornia,
Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and
Superscript I RT (Life Technologies). Note: this is a
NIH\_MGC Library."
a 224 c 204 g 116 t Site\_2: lly cloned

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US-09-993-234-6
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Soares_fetal_heart_NbHH19W Homo sapiens cDNA clone
710 3' similar to SW:WSL1_HUMAN Q93038 WSL-1 PROTEIN
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LeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGlyTrpPheVal
                                                                H1sH1sAsnSerGluCysAlaArgCysGlnAlaCysAspGluGlnAlaSerGlnValAla
                                                                                                                                                                                                       Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.lngov) for further information.
Insert Length: 1640 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 494.
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Contact: Robert Strausberg, Ph.D
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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/db_xref="taxon:9606"
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GluCysGlnValSerGlnCysValSerSerSerProPheTyrCysGlnProCysLeuAsp 143
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: yongsung@mail.kribb.re.kr
Plate: 48 row: G column: 11
High quality sequence stop: 647.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Korea Research Institute of Bioscience & Biotechnology 52 Eceun dong Yuseong-gu, Daejeon 305-333, South Korea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (2002)
Contact: Kim YS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tel: +82-42-860-4470
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Kim, N.S., Hahn, Y.,
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/note="Organ: Stomach; Vector: pT773-Pac; Site_1: EcoRI; Site_2: NotI; The $22$NU16 library was contributed by the Soares laboratory and it was constructed as described by Bonaldo, M.F., Lennon, G. and Soares, M.B. (1996), Genome Research 6(9): 791-806. RNA was prepared from harvested cells of $NU-16 culture. $NU-16 cell was obtained from Korean Cell Line Bank (KCLB). $NU-16 was established from ascitic fluids of Korean patients by Park J.G. et al. (1990), Cancer Res 50: 2773-2780."
                                                                                                                                                                                                                                      /tissue_type="Ascites"
/cell_type="Lymphoblast-like"
/cell_line="SNT_16"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
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Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
Proceeding the consortium of the contact the contact the contact the contact info@image.llnl.gov) for further information.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 585)
1 (bases 1 to 585)
1 (Canpanta Cancer Institute, Cancer Genome Anatomy Prof
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AW517358
AW517358.1 GI:7155440
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        LeuGlySerCysProGluArgCysAlaAlaValCysGlyTrpArgGln 199
                              CTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCCAGGCTGGTTTGTG
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Location from Gibco
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Plate: 31 row: H column: 06
High quality sequence stop: 552.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Korea Research Institute of Bioscience & Biotechnology 52 Eceun-dong Yuseong-gu, Daejeon 305-333, South Korea Tel: +82-42-860-4470 Fax: +82-42-860-4409
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kim,N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim,M.R.,Oh,K.J., Cheong,J.E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,S. and
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                                                                                                                                                                                                                                                                                                                                                                            //note-Togans Stomach; Vector: pCNS; Site_1: EcoRI; Site_2: NotI; The poly (A)+ RNA was dephosphorylated with bacterial alkaline phosphatase (BAP) and then decapped with tabacco acid pyrophosphatase (TAP). The decapped intact mRNA was ligated with DNA-RNA linker including EcoR I site by treatment of T4 RNA ligase and the first strand cDNA was synthesized from oligo dT-selected mRNA by priming with dT-tailed vector. The dT-tailed vector was adjusted to have about 60nt. The cDNA vector was circularized with E. coli DNA ligase after digestion of EcoRI which site is also included in vector. An RNA strand conserved to a DNA strand by Okayama-Berg method. The obtained cDNA vectors were used for transformation of competent cells E. coli ToplOF' by electroporation method. The cDNA libraries constructed by this method are full-length enriched cDNA library."
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/cell_line="SNU-620"
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/db_xref="taxon:9606"
/clone="S6SNU620-31-H06"
/clone_lib="S6SNU620"
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                              Email: yongsung@mail.kribb.re.kr
Plate: 17 row: C column: 06
                                                                                                                                                                                                                                                                                Korea Research Institute of Bioscience & Biotechnology 52 Boeun-dong Yuseong-gu, Daejeon 305-333, South Korea Tel: +82-42-860-4470
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BM744307
BM744307.1 GI:19065636
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lm, N.S., Hahn, Y.,
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1 to 565)
/note="Organ: Stomach; Vector: pCNS; Site_1: EcoRI;
Site_2: NotI; The poly (A)+ RNA was dephosphorylate
                                                                                                                              /organism="Homo sapiens"
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/clone="S6SNU620-17-C06"
                                                 /cell_type="Scattering floating"
/cell_line="SNU-620"
                                                                              /tissue_type="Ascites"
                                                                                                 /sex="F"
                 /note="Organ:
                                  /lab_host="Top10F'"
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Plate: 27 row: G column: 03
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Kim, M.S., Hahn, Y., Oh, J.H., Lee, J.Y., Ahn, H.Y., Chu, M.Y., Kim, M.R., Oh, K.J., Cheong, J.E., Sohn, H.Y., Kim, J.M., Park, H.S., Kim, S. and
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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/cell_line="SNU-620"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kim,N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y.,
Oh,K.J., Cheong,J.E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,
Xim,Y.S.
21C Frontier Korean EST Project 2001
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1 (bases 1 to 514)

Kim, N.S., Hahn, Y., Oh, J.H., Lee, J.Y., Ahn, H.Y., Chu, M.Y., Kim, M.R
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                                                                                                                                                                                                                                                                                                                       Email: yongsung@mail.kribb.re.kr
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        /note-*Organ: Stomach; Vector: pCNS; Site_1: EcoRI; Site_2: NotI; The poly (A)+ RNA was dephosphorylated with bacterial alkaline phosphatase (BAP) and then decapped with tabacco acid pyrophosphatase (TAP). The decapped intact mRNA was ligated with DNA-RNA linker including EcoR I site by treatment of T4 RNA ligase and the first strand cDNA was synthesized from oligo dT-selected mRNA by priming with dT-tailed vector. The dT-tailed vector was adjusted to have about 60nt. The cDNA vector was circularized with E. coli DNA ligase after digestion of
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converted to a DNA strand by Okayama-Berg method. To obtained cDNA vectors were used for transformation competent cells E. coli ToplOF' by electroporation The cDNA libraries constructed by this method are full-length enriched cDNA library." method.

Percent Similarity:
Best Local Similarity:
Query Match: US-09-993-234-6 (1-417) x BM783979 14 AlaLeuLeuLeuValLeuClyAlaArgAlaGlnGlyGlyThrArgSerProArgCys LeuCysSerArgArgAspThrAspCysGlyThrCysLeuProGlyPheTyrGluH1sGly 173 GlyHisTyrLeuLysAlaProCysThrGluProCysGlyAsnSerThrCysLeuValCys SerProPheTyrCysGlnProCysLeuAspCysGlyAlaLeuHisArgHisThrArgLeu 153 ArgCysGlyCysLysProGlyTrpPheValGluCysGlnValSerGlnCysValSerSer AspGlyCysValSerCysProThr CGCTGTGGCTGTAAGCCAGGCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGT GGGCACTACCTGAAGGCCCCTTGCACGGAGCCCTGCGGCAACTCCACCTGCCTTGTGTGT AspCysAlaGlyAspPheHisLysLysIleGlyLeuPheCysCysArgGlyCysProAla GCGCTCCTCCTGGTGCTGCGGGGCCCCGGGCCCCAGGGCGCACTCGTAGCCCCAGGTGT TCACCCTTCTACTGCCAACCATGCCTAGACTGCGGGGCCCTGCACCGCCACACACGGCTA Tumor Gene Index Unpublished (1997) NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap National Cancer Institute, Cancer Genome Ana Eukaryota; Metazoa; Mammalia; Eutheria; 1 (bases 1 to 508) EST PRECURSOR ;, mRNA sequence. AI203624 AI203624 508 bp mRNA linear EST 29-OCT-qe75f02.xl Soares\_fetal\_lung\_NbHL19W Homo sapiens cDNA clone IMAGE:1744827 3′ similar to SW:WSL1\_HUMAN Q93038 WSL-1 PROTEIN Homo sapiens AI203624.1 86 Robert 1.19e-72 984.00 100.00% 100.00% 42.36% GI:3756230 Strausberg, Chordata; Primates; 181 508 (1-514)508 bp Length:
Matches:
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Mismatches:
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This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
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Search completed: April 6, 2003, 23:25:07 Job time : 2620.38 secs

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## ALIGNMENTS

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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1
CURRENT APPLICATION NUMBER: US/08/815 450
FILING DATE: HERFUTF...
CLASSIETT.
                         PRIOR APPLICATION NUMBER: NO. 61
APPLICATION NUMBER: NO. 61
PETLING DATE: 06-FEB-1997
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 60/
FILING DATE: 17-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
              PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Death Domain Containing Receptors NUMBER OF SEQUENCES: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                         STREET: 1100 New CITY: Washington
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
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1100 New York Ave.,
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Gentz, Reiner L.
Dillon, Patrick
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Ni, Jian
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US 60/013,285
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Best Local Similarity:
Query Match:
DB:
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INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 1254 base pairs
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STRANDEDNESS: doub
TOPOLOGY: both
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REGISTRATION NUMBER: 36,
REFERENCE/DOCKET NUMBER:
CAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCCAGGC
                                                                                                                                      TGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCCTTCTACTGCCAACCA
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Indels:
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Matches:
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                              COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb flo
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WINPATIN (Genentech)
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/928,069
FILING DATE: 11-Sep-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: 60/026943
PRIOR APPLICATION UMBER: 60/026943
FILING DATE: 109/23/1996
ATTORNEY/AGENT INFORMATION:
NAME: MAISCHARP, Diane L.
BEGITERATION:
                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                    APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-3 PC
NUMBER OF SEQUENCES: 15
NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: 910
TELECOMMUNICATION INFORMATION:
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CITY: S
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California
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TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity:
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LENGTH: 1634 base pairs
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STRANDEDNESS: Single
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                                               CysLeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThr
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                         MetGluAlaLeuThrProProProAlaThrHisLeuSerProLeuAspSerAlaHisThr
                                                                                                                                                                                   TGCCTAGACTGCGGGGCCCTGCACCGCCACACGGCTACTCTGTTCCCGCAGAGATACT
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Conservative:
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Patent No. 6469144
GENERAL INFORMATION:
APPLICANT: AShKenazi, Avi J.
TITLE OF INVENTION: Apo-2 LI AND
NUMBER OF SEQUENCES: 28
                          TELEPHONE: 650/225-5416
TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 1634 base pairs
TYPE: Nucleic Acid
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                                                                                                          REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007P1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/828,683A
FILING DATE: 31-Mar-1997
CLASSIFFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 i
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                                                                                                                                                                               APPLICATION NUMBER: 08/625328
FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech)
                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: South San
           STRANDEDNESS: Single
                                                                                                                                                    NAME: Marschang, Diane L.
                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
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TOPOLOGY:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, \
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/815,469
FILING DATE: HEREWITH
CLASSIFICATION LORIA:
APPLICATION DATA:
APPLICATION NUMBER: NO. 6153402 Y
FILING DATE: 06-FEB-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 1, Application Patent No. 6153402
                                     TELEFAX: 202-371-2540
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                  REFERENCE/DOCKET NUMBER: 14
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
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APPLICANT:
                                                                                                                             APPLICATION NUMBER: US 60
FILING DATE: 12-MAR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Steffe, Eric K.
REGISTRATION NUMBER: 36,0
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APPLICATION NUMBER: US 6(
FILING DATE: 17-OCT-1996
PRIOR APPLICATION DATA:
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STREET: 1100 New York Ave., N
CITY: Washington
STATE: DC
COUNTRY: USA
ZIP: 20005-3934
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APPLICANT: Gentz, Reiner L.
APPLICANT: Dillon, Patrick
TITLE OF INVENTION: Death D
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Ni, Jian
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Indels:
Gaps:
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SULT 5 -08-928-069-5 -08-928-069-5 -08-928-069-5 Sequence 5, Appl Patent No. 64622 GENERAL INFORM APPLICANT: FITTLE OF INVI NUMBER OF SEC CORRESPONDEN ADDRESSEE: 1 CITY: SOUT STATE: Call COMPUTER REAL MEDIUM TYPE COMPUTER REAL MEDIUM TYPE COMPUTER: OPERATING S SOFTWARE: OPERATING DATE CLASSIFICATION FILING DATE APPLICATION FILEPHAN: MARE: REFERENCE,/I TELEPHONE: TELEPHO								
95 9069-5 907-5 908-6 90	P Qy	ОУ	D Qy	Оy	D Q	Db	A d	Qy dd
5, Appl 5, Appl 1, App	417 1479	397 1419	377 1359	357 1299	337 1239	317 1179	297 1119	277 1059
TION: ISAkemazi, Avi J. ISAkemazi, Avi J. ISAkemazi, Avi J. ISA ADO-3 POLYPEPTIDE  DUENCES: 15 E ADDRESS: Genentech, Inc. DNA Way IF San Francisco Lifornia ISA  DABLE FORM: ISA  ISA  ISA  ISA  ISA  ISA  ISA  ISA	Pro 417      CCG 1481	LeuGluargMetGlyLeuAspGlyCysValGluAspLeuArgSerArgLeuGlnArgGly 416 	GluMetLeuLysArgTrpArgGlnGlnGlnProAlaGlyLeuGlyAlaValTyrAlaAla 396 	ArgGluAlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyr 376	AspValMetAspAlaValProAlaArgArgTrpLysGluPheValArgThrLeuGlyLeu 356 	SerProGluSerProAlaGlySerProAlaMetMetLeuGlnProGlyProGlnLeuTyr 336	ThrTrpSerTrpAspGlnLeuProSerArgAlaLeuGlyProAlaAlaAlaProThrLeu 316	ValGlyAsnSerTrpThrProGlyTyrProGluThrGlnGluAlaLeuCysProGlnVal 296 

Score:

Alignment Scores:

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Query Match:

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Percent Similarity:
Best Local Similarity:
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                                                                                                                                                                                        equence 5, Application US/08828683A atent No. 6469144
GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-2 LI ANDWAREN OF SEQUENCES: 28
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                        COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WilPatin (Genentech)
                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                  ThrSerThrLeuGlySerCysProGluArgCysAlaAlaValCysGlyTrpArgGln 199
                                                                                                                                                                                                                                                                                                                                                                ACGTAATTCCTA---
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STREET: 1 DNA WAY
CITY: South San Francisco
STATE: California
COUNTRY: USA
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Query Match:
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Pred. No.:
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REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P10
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650,7225-5416
                   ThrSerThrLeuGlySerCysProGluArgCysAlaAlaValCysGlyTrpArgGln 199
                                                                        AspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysPro 180
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                                                                                                                                                                                TrpPheValGluCysGlnValSerGlnCysValSerSerSerProPheTyrCysGlnPro
                                                        GACTGTGGGACCTGCCTGCCTGCCTTCTATGAACATGGCGATGGCTGCGTGTCCTGCCCC
                                                                                                              TGCCTAGACTGCGGGGCCCTGCACCGCCACACACGGCTACTCTGTTCCCGCAGAGATACT
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TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
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APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
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US-08-928-069-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Marschang, Diane L.
REGISTON NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P10
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
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NAME: Marschang, Diane L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Apo-3 POLYPEPTIDE NUMBER OF SEQUENCES: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                         No . .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 433 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
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FILING DATE: 09/23/1996
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FILING DATE: 11-Sep-1997
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SOFTWARE: WinPati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY:
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sProGlyTrpPheValGluCysGln-ValSerGlnCysValSerSer-SerProPheTyr 137
                                                                                          sAlaProCysThrGluProCysGlyAsnSerThrCysLeuValCysProGlnAspThrPh 78
                                                                                                                                                                                                                                   PheHisLysLysIleGlyLeuPheCysCysArgGlyCysProAla-GlyHisTyrLeuLy
                                                                                                                                                                                                                                                                                CTGCTGGGGGCCCAGGGCGCAGGCGCACTCGTAGCCCCCAGGTGTGACTGTGCCGGTGAC
                             GGCCTCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCCGCTGTGGCTGTAA
                                             nAlaSerGlnValAlaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLy 118
                                                                                                                                                      GGCCCCTTGCACGGAGCCCTGCGC-AACTCCACCTGCCTTGTGTGTCCCCCAAGACACCTT
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SYSTEM: PC-DOS/MS-DOS
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                                                              1 CTGCTGGGGGCCCGGGCCAGNGGCGGCACTCGTAGCCCCAGGTGTGACTGTGCCGGTGAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: 3.5 inch, 1.44 mb floppy COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech) CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/625328
FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-5ep-1996
                                                                                                                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
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TELEPHONE: 650/225-5416
TELEFAX: 650/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-2 LI AND Apo-3 POLYPEPTIDES
NUMBER OF SEQUENCES: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                 LeuLeuGlyAlaArgAlaGlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAsp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gArgAsp 159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TTCCACAAGAAGATTGGTCTGTTTTGTTGCAGAGGCTGCCCAGCGGGGCAACTACCTGAA 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TGCCAACCATGCCTAGACTGCGGGGCCCTGCAACGCCAACACGCCTAATNTGTTTCCCG 419
                                                                                                                                                                                                                                                                                                                                                  TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/828,683A FILING DATE: 31-Mar-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Genent
STREET: 1 DNA Way
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: P1007P1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Marschang, Diane L. REGISTRATION NUMBER: 35,600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: South San
                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 433 base pairs
                                                                                                                                                                                                                                                                                                                   DESCRIPTION: SEQ ID
                                                                                                                                                                               1.74e-48
692.00
93.10%
91.72%
29.79%
                                                                                                                                                                                                                                                                                                                                    Linear
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Matches:
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133
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Percent Similarity:
                              Pred. No.:
                                           Alignment Scores:
                                                                        US-08-762-308-10
                                                                                                                                                        APPLICANT: Beller, Bruce A.
APPLICANT: Bezoni, Flavia M.
TITLE OF INVENTION: MODIFIED RECEPTORS THAT CONTINUOUSLY
TITLE OF INVENTION: SIGNAL
CORRESSOR ADDRESS:
ADDRESSEE ATOOLG, White & Durkee
STREET: P.O. Box 4433
CITY: Houston
STATE: TX
COUNTRY: USA
ZIP: 77210-4433
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/08/762,308
FILING DATE: 09-DEC-1996
CLASSIFICATION NUMBER: US/08/762,308
FILING DATE: 09-DEC-1996
CLASSIFICATION NUMBER: US/08/762,308
FILING DATE: 05-APR-1994
ATTOONEY/AGENT INFORMATION:
NAME: Kitchell, Barbara S.
REGISTRATION NUMBER: UTSD: 335--1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 418-3000
TELEPHONE: 418-3000
TELEPHONE: 418-3010
INFORMATION FOR SEO ID NO: 10:
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                                                                                                                     TELEFAX: 474-7577
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 1956 base pairs
TYPE: TIPLE AND A SAIR
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                                                                                    STRANDEDNESS:
TOPOLOGY: 11
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                                                                                                            nucleic acid
                                                                                     linear
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39.19%
Length:
Matches:
Conservative:
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Qy 282ThrProGlyTyrProGluThrGlnGlnAlaLeuCys	271 987	258 927	238 867	807	205 747	687	642AGCCACTGCAAGAAAAAT	180 ProThrSerThrLeu	Qy 160 ThraspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCys	528 GACTGCAGCCCCTGC	480 - 140 P	120 GlyTrpPheVa	Qy 100 SerGlnValAlaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysPro	80 AlaTrpGluAsnHisHisAsnSe     :::   :: 366 GCTTCCCAGAATTACCTCAGG	Qy 60 ProCysThrGluProCysGlyAsnSerThrCysLeuValCysProGlnAspThrPheLeu	· N	186 GTCCCTTCTCTTGGTGACCGGGAGAAG	ш		Hest Local Similarity: 27.198 Mismatches Ouery Match: 17.418 Indels: DB: 2 Gaps:
JeuCys	VIGCCACCTGTCAGTGAGGTA 1043	ProProAspSerSerGluLys 270            CTCCTGTCTCCAGTACCCCC 986	HisLeuSerProLeuAspSer 257         CCTTCAGCCCCACCTCCGGC 926	LysProLeuValThrAlaAsp 237     ::: 	AlaThrLeuThrTyrThrTyr 224 :::    ATCAGTTTAATGTGCCGATAT 806	TCAGGTACTGCGGTGCTGTTGCCCCTG 746		GluArgCysAlaAlaValCys 195	GlyaspGlyCysValSerCys 179         GAAAGTGAGTGCGTCCCTTGC 641	-TTCAACGGCACCGTGACAATCCCCTGTAAGGAGACTCAG 581	หลั	SerSerProPheTyrCysGln 139	ThrargCysGlyCysLysPro 119               ACGGTGTGTGGCTGTAAG 479	rGlucysAlaArgCysGlnAlaCysAspGluGlnAla 99 :::	CysProGlnAspThrPheLeu 79     :::         !TGTGAAAAGGGCACCTTTACG 365	AlaGlyHisTyrLeuLysAla 59            ::: AAAGGAACCTACTTGGTGAGT 305	ArgSerProArgCysAspCysAlaGlyAspPhe 39     	yAlaGlnGlyGlyThr 28          CCATCAGGGGTCACTGGACTA 185	56)	s: 191 93 18

Db 333 GTGCTTCCAGCTCTGTTGGCAGATGTGTACCCCGCAGGGGTTCAGGGGCTGGTCCCTCAC 392  Qy 31 ProArgCysAspCysAlaGlyAspPhe39	-09-993-234-6 (1-417) x US-09-513-007-1 (1-2440)  11 ValalaalaalaLeuLeuValLeuLeuGlyAlaargalaGlnGlyGlyThrArgSer	Match: 16.68% Indels: 4 Gaps:	Pred. No.: 1.98e-22 Length: 2440 Score: 387.50 Matches: 130 Percent Similarity: 36.57% Conservative: 47 Best Local Similarity: 26.86% Mismatches: 186	; LOCATION: (294)(1706) US-09-513-007-1 Alignment Scores:	TYPE: DNA ORGANISM: FEATURE: NAME/KEY:	SOFTWARE: FastSEQ for Windows Version 4.0  SEQ ID NO 1  LENGTH: 2440	CURRENT APPLIC CURRENT FILING PRIOR APPLICAT PRIOR FILING D	; APPLICANT: MWANG1, SIMON ; TITLE OF INVENTION: BOVINE TUMOR NECROSIS FACTOR RECEPTOR-1 ; TITLE OF INVENTION: AND METHODS OF USE ; FILE REFERENCE: 08411-018001	GENERAL INI APPLICANT APPLICANT APPLICANT	RESULT 10 US-09-513-007-1 ; Sequence 1, Application US/09513007 ; Patent No. 6406907	Qy 411 SerArgLeuGlnArgGlyPro 417	Qy 391 GlyAlaValTyrAlaAlaLeuGluArgMetGlyLeuAspGlyCysValGluAspLeuArg 410	Qy 374 GinGinTyrGiuMetLeuLysArgTrpArgGinGinProAlaGlyLeu 390	Oy 355 GlyLeuArgGluAlaGluIleGluAlaValGluValGluIleGlyArgPheArgAsp 373	Qy 335 LeuTyrAspValMetAspAlaValProAlaArgArgTrpLysGluPheValArgThrLeu 354	Qy 315 ThrLeuSerProGluSerProAlaGlySerProAlaMetMetLeuGlnProGlyProGln 334 :::    :::     1128 TCCGCCCACCCGCAACGTCCTGACAATGCAGACCTTGCGATT	Qy 295 GlnValThrTrpSerTrpAspGlnLeuProSerArgAlaLeuGlyProAlaAlaAlaPro 314  Db 1104 ACCTCTGTTCAGAAATGGGAAGAC:
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Qy	Оу	Qу	Db Qy	Ор	Db Qy	Db Cy	Qу	Дb	Qу	Qу	Оу	Db Qy	Дb	Дb	Db Qy	Qу	Оy
314 ProThrLeuSerProGluSerProAlaGlySerProAlaMetMetLeuGlnProGlyPro    :::        1386 GATCAGCTCGCGGATGCCGACCCCGCG		276 LeuValGlyAsnSerTrpThrProGlyTyrProGluThrGlnGluAlaLeuCys :::::	256 AspSerAlaHisThrLeuLeuAlaProProAspSerSerGluLysIleCysThrValGln	236 AlaAspGluAlaGlyMetGluAlaLeuThrProProProAlaThrHisLeuSerProLeu	232ProLeuValThr	231 1047 CTGGTAAAAGAGGGGAGCCAGAACTCCTGGTCCCGGCCCCAGGCTTCAACCCCACCACC	222 TyrThrTyrArgHisCysTrpProHisLys231	202 TrpValGlnValLeuLeuAlaGlyLeuValValProLeuLeuLeuGlyAlaThrLeuThr 221    ::::       ::: 930 TTACCCCTGGTGATTGTCTTCGGGCTTTGCCTGGCATCCTTCGCCTCTGTCGTCTTAGCA 989	194 ValCysGlyTrpArg	175 GlyCysValSerCysProThrSerThrLeuGlySerCysProGluArgCysAlaAla 193	156SerargargaspThraspCysGlyThrCysLeuProGlyPheTyrGluH1sGlyAsp 174	138 CysGlnProCysLeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCys 155	118 LysProGlyTrpPheValGluCysGlnValSerGlnCysValSerSerSerProPheTyr::::	98 GlnalaSerGlnValalaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCys :::         ::     615 GAAATGTTCCAGGTGGAGATTTCGCCTTGTGTAGTGGACCGGGACACTGTGTGCGGCTGC	78 PheLeuAlaTrpGluAsnHisHisAsnSerGluCysAlaArgCysGlnAlaCysAspGlu:::	58 LysalaproCysThrGluProCysGlyAsnSerThrCysLeuValCysProGlnAspThr	40HislyslyslleglyLeuPheCysCysArgGlyCysProAlaGlyHisTyrLeu     :::

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Percent Similarity:
Best Local Similarity:
Query Match:
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US-08-050-319B-24
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                                                                                      Score:
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                                                                                                                        Alignment Scores:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                               TELEFAX: (415) 327-323 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                NAME: Robbins, Roberta L.
REGISTRATION NUMBER: 33,208
REFERENCE/DOCKET NUMBER: 51
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 617-8999
                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 10-May-1993
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: M.Feldmann, P.W. Gray,
APPLICANT: M.J.C. Turner, F.M Brennan
TITLE OF INVENTION: Modified human TNFalpha (Tumor
TITLE OF INVENTION: Necrosis Factor alpha) Receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1473
                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1653 GAGGAGGCGCTG 1664
                                                                                                                                                                                                                              MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                              TOPOLOGY: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: FILING DATE: 10-May
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
                                                                                                                                                                                            NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                              LOCATION:
                                                                                                                                                                                                                                                                                               LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 T: 635 Bryant Street Palo Alto
                                                                                                                                                                                                                                                                               H: 2062 base pairs nucleic acid
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327-3231
NO: 24:
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39.09%
28.29%
16.19%
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                                                                                    Length: Matches:
                                                  Conservative: Mismatches:
                                     Indels:
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131
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CAGGGGGCTGACCCCATCCTTGCGACAGCCCTCGCCTCCGACCCCATCCCCAACCCCCTT 1168
                                                                                               CysProGlnValThrTrpSerTrpAspGlnLeuProSerArgAlaLeuGlyPro-----
                                                                                                                                                                                                       CCCACTCCAGGCTTCACCCCCACCCTGGGCTTCAGTCCCGTGCCCAGTTCCACCTTCACC 1036
                                                                                                                                                                                                                                                                                                            GlyMetGluAlaLeuThrProProProAlaThrHisLeuSerProLeuAspSer-----
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                                                                                                                                      TCC-----AGCTCCACCTATACCCCCGGTGAC------
                                                                                                                                                          ThrValGlnLeuValGlyAsnSerTrpThrProGlyTyrProGluThrGlnGluAlaLeu 292
                                                                                                                                                                                                                                                                                                                                              ---TGGAAGTCCAAGCTCTACTCCATTGTTTGTGGGAAATCGACACCTGAAAAAGAGGGG
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                                                                TGTCCCAACTTTGCG------GCTCCCCGCAGAGAGGTGGCACCACCCTAT 1108
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                                -----AlaAlaAlaProThrLeuSerPro---
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US-08-465-982-24
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US-08-465-982-24
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Patent No. 5863786
GENERAL INFORMATION:
APPLICANT: M.Feldmann, P.W. Gray,
APPLICANT: M.J.C. Turner, F.M Brennan
TITLE OF INVENTION: Modified human TNFalpha (Tumor
TITLE OF INVENTION: Necrosis Factor alpha) Recepton
                                                                                                                      TELEFAX: (415) 327-323
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 2062 base pair:
              FEATURE:
NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08
FILING DATE: 10-May-1993
ATTORNEY/AGENT INFORMATION:
NAME: Robbins, Roberta L.
                                                                                                                                                                                REGISTRATION NUMBER: 33,208
REFERENCE/DOCKET NUMBER: 51:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 617-8999
                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                            MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
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TCC----
                           ThrValGlnLeuValGlyAsnSerTrpThrProGlyTyrProGluThrGlnGluAlaLeu
                                                          CCCACTCCAGGCTTCACCCCCACCCTGGGCTTCAGTCCCGTGCCCAGTTCCACCTTCACC
                                                                                                                        GAGCTTGAAGGAACTACTACTAAGCCC------CTGGCCCCAAACCCAAGCTTCAGT
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US-09-106-038A-1
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GENERAL INFORMATION:
                                                                                                                  TELEFAX: (760) 603-382
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                       TELEPHONE: (760) 931-9200
                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Laurel Spear Bernstein
REGISTRATION NUMBER: 97,280
REFERENCE/DOCKET NUMBER: RTS-
                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible OPERATING SYSTEM: Windows NT SOFTWARE: Microsoft Word 97 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch disk, 1.44 Mb
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                                                                                                                                                                                                                                                                             FILING DATE: JUCCLASSIFICATION:
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                                                                                                                 US-09-993-234-6 (1-417) x US-09-505-250-3 (1-2161)
                                                                                                                                                                          Percent Similarity:
Best Local Similarity:
                                                                                                                                                                                                                                     Alignment Scores:
                                                                                                                                                              Query Match:
                                                                                                                                                                                                                        Pred.
                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Fas
SEQ ID NO 3
LENGTH: 2161
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 3, Application US/09505250A Patent No. 6329148
                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Rosen, Glenn
APPLICANT: Kao, Peter
TITLE OF INVENTION: Synergistic Anti-Cancer Therapy with
TITLE OF INVENTION: Triptolides and Death Domain Ligands
FILE REFERENCE: SUN-109PRV2
                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/0 CURRENT FILING DATE: 2000-02-15
                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                        ORGANISM: H. sapiens
                                                                                                                                                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
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GGGGACAGGGAGAAGAGAGATAGTGTGTGTCCCCAAGGAAAATATATCCACCCTCAAAAT
                          GlyThrArgSerProArg------CysAspCysAlaGlyAspPheHisLysLysIle
                                                       CTCCTGGAGCTGTTGGTGGGAATATACCCCTCAGGGGTTATTGGACTGGTCCCTCACCTA
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                                                               CAGAAGTGGGAGACAGCGCCCACAAGCCACAGAGCCTAGACACTGATGACCCCGCGACG
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US-08-321-668-1
                          Percent Similarity:
Best Local Similarity:
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US-08-321-668-1
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                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: WA
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
                                                                             No.:
                                                                                                                                                                             FEATURE:
                                                                                                                                                                                           TOPOLOGY: 1:
MOLECULE TYPE:
                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 12-OCT-1993
ATTORNEY/AGENT INFORMATION:
NAME: BROWDY, ROGER L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: WALLACH, David
APPLICANT: BARKEBUSCH, COrd
APPLICANT: VARFOLOMEEV, Eugene
APPLICANT: BATKIN, Michael
TITLE OF INVENTION: THE THE RECEPTORS, THEIR PREPARATIO
             Match:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1510
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TELEFAX: 248633
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                                                                                                                                                            NAME/KEY:
                                                                                                                                                                                                                                                                                                        TELEPHONE: 202-737-3528
                                                                                                                                                                                                                                                                                                                                                                                    NAME: BROWDY, Roger L. REGISTRATION NUMBER: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/0 FILING DATE: 12-OCT-1994
                                                                                                                                                                                                                           STRANDEDNESS:
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             411 SerArgLeu 413
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Search completed: April 6, 2003, 23:29:19 Job time: 96.3944 secs

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Command line parameters:

-MODEL-frame+_p2n.model -DEV=xlh
-Q-/cgn2_1/USPT0_spool_VS09993234/runat_27032003_115455_15349/app_guery.fasta_1.2346
-Q-/cgn2_1/USPT0_spool_VS09993234/runat_27032003_115455_15349/app_guery.fasta_1.2346
-DB-GenEmbl -QFMT-fastap -SUFFTX-rge -MINMATCH-0.1 -LOOPCL-0 -LOOPEXT-0
-UNITS-bits -START-1 -END--1 -MATRIX-blosum62 -TRANS-human40.cdi -LIST-45
-DCCALIGN-200 -THR_SCORE-pct -THR_MAX-100 -THR_MIN-0 -ALIGN-15 -MODE-LOCAL
-DUTFMT-pto -NORM-ext -HEAPSIZE-500 -MINLEN-0 -MAXIEN-20000000000
-USER-US0993234_@CCN_1_1_8534_@runat_27032003_115455_15349 -NCPU-6 -ICPU-3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES-0 -WAIT -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=90 -THREADS-1 -XCAPOP=10 -XCAPEXT-0.5 -FGAPOP=6 -FGAPEXT=7
-YGAPOP=10 -YGAPEXT-0.5 -DELOP=6 -DELEXT-7
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Submitted (19-MAR-1997) Molecular Immunology Group, Institute of
Molecular Medicine, John Radcliffe Hospital, Headington, Oxford
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1 (bases 1 to 1143)
Screaton,G.R., Xu,X.N., Olsen,A.L., Cowper,A.E., Tan,R.,
McMichael,A.J. and Bell,J.I.
LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing
Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
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/db_xref="GI:2071967"
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GluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGlyTrpPheValGlu
                  TGCGGCAACTCCACCTGCCTTGTGTGTCCCCAAGACACCTTCTTGGCCTGGGAGAACCAC
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                                                                                              CTGTTTTGTTGCAGAGGCTGCCCAGCGGGGCACTACCTGAAGGCCCCTTGCACGGAGCCC
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                                                                                                                                                                                                                                                                                                                     1 (bases 1 to 1254)
Yu,G.-L., Ni,J., Gentz,R.L. and Dillon,P.J.
Death domain containing receptors
Patent: US 6153402-A 3 28-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
AR119657
                                                                                                                                                                                                                                                                                                                                                                      Unknown
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1 (bases 1 to 1254)
Chinnaiyan, A.M., O'Rourke, K., Yu, G.L., Lyons, R.H., Garg, M., Duan, D.R., Xing, L., Gentz, R., Ni, J. and Dixit, V.M.

Signal transduction by DR3, a death domain-containing receptor related to TNFR-1 and CD95
Science 274 (5289), 990-992 (1996) NOV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chinnaiyan, A.M., O'Rourke, K., Yu, G.-L., Lyons, R.H., Duan, R., Xing, L., Gentz, R., Ni, J. and Dixit, V.M. Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human death receptor U72763
                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (29-SEP-1996) Pathology, University of Michigan Medical School, 1301 Catherine St., Box 0602, Ann Arbor, MI 48109, USA
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                                                                                                                         201
                                                                                                                                                              /product="death receptor 3"
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TLTYTYRHCWPHKPLVTADEAGMEALTPPPATHLSPLDSAHTLLAPPDSSEKICTVQL
VGNSWTPGYTETGEALCPQVWSWDQLLPSRALGPAAAPTLSPESPAGSPAMILQPGPQ
                                                                                                                       LYDVMDAVPARRWKEFYRTLGLREDE EAVEVEIGRFRDQOYEMLKRWRQQQPAGLGA
VYAALERWGLDGCVEDLGRLGRGP E
1 420 c 407 g 226 t
                                                                                                                                                                                                                                                                                                                /gene="DR3"
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/db_xref="taxon:9606"
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Submitted (12-NOV-1996) Blochemistry,
WA 98101, USA
                                                                                                                                                                                                                                        of Apoptosis
Unpublished
                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 1254)
                                                                                                                                                                                                            Degli-Esposti, M.A. and Goodwin, R.G.
                                                                                                                                                                                                                                                                                                  Degli-Esposti, M.A.,
                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
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                                                                                                                                                                                                                                                                    A Novel Member of the TNF Receptor Family, Is a Strong Inducer
                                                                                                                                                                                                                                                                                                                                                                                                                                    aoptosis inducing
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                 receptor AIR*
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Query Match:
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υ94501.1
    Screaton, G.R., Xu, X.N., Olsen, A.L., McMichael, A.J. and Bell, J.I.
                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 1254)
                                                                                                                                            \begin{array}{lll} {\rm HSU}94501 & {\rm 1254~bp} & {\rm mRNA} \\ {\rm Human~lymphocyte~associated~receptor~of} \end{array}
                                                                      Homo
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VYAALERMGLDGCVEDLRSRLQRGP"
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                                                       TGCGGCAACTCCACCTTGTGTGTGTCCCCAAGACACCTTCTTGGCCTGGGAGAACCAC
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/protein_id="AAC51306.1"
/db_xref="G1:2071949"
/db_xref="G1:2071949"
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DCGTCLDGFYEHGDGCVSCPTSTLGSCDERCAAVCGWROMFWVOVLLAGLVVPLLIGA
TLTYTYRHCWPHKPLVTADEAGMEALTPPPATHLSPLDSAHTLLAPPDSSEKICTVQL
VGNSWTPGYPETGCALCPQVTWSWDQLDSRALGPAAAPFLSPESPAGSPAMMLQPGPQ
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/db_xref="taxon:9606"
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/note="LARD-1a; membrane prote
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Submitted (19-MAR-1997) Molecular Immunology Group, Institute of Submitted (19-MAR-1997) Molecular Immunology Group, Oxford
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LARD: a new lymphoid-specific death domain containing regulated by alternative pre-mRNA splicing Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
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                                                                                                                                                      /note="insertion compared to LARD-1a, deposited in GenBank Accession Number U94501, probably represents alternative 3' splice site"
1 421 c 408 g 226 t
                                                                                                                                                                                                                                                                /product="lymphocyte associated receptor of death 1b"
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                                                                                                                                                                                                                                                                                                                                                                                                /function="mediates apoptosis"
/note="LARD-1b; membrane prote
TNF-R1; contains a death domai
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LARD: a new lymphoid-specific death domain containing re regulated by alternative pre-mRNA splitcing
Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
                                                                                                                                                                                                                                                           Submitted (19-MAR-1997) Molecular Immunology Molecular Medicine, John Radcliffe Hospital,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/function="mediates apoptosis"
/note="LARD-2; similar to Fas and TNF-R1; postuble; truncated before the transmembrane
                                                                                                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                         /cell_line="HeLa"
                                                                                                                                                                                                                Location/Qualifiers
                                                                                              _type="lymphocyte"
                                                                                                                                                                                                                                                           Group, Institute of Headington, Oxford OX3
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                     GI:1763292
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Accession Number U94501, probable retained intron leading
to premature transcriptional termination"
441 c 451 g 240 t
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/product="lymphocyte associated receptor
/protein_id="AAC51308.1"
/db_xref="GI:2071953"
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Marsters, S.A., Sheridan, J.P., Donahue, C.J., Pitti
Goddard, A.D., Bauer, K.D. and Ashkenazi, A.
Apo-3, a new member of the tumor necrosis factor
contains a death domain and activates apoptosis a
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Goddard, A.D., Bauer, K.D.
Direct Submission
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/dev_stage="fetal"
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NF-kB; TNF receptor family member"
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1 (bases 1 to 1652)
Ashkenazi,A.J., Baker,K.P., Chan,B., Goddard,A., Godowski,P.J., Gurney,A.L., Hebert,C., Henzel,W., Kabakoff,R.C., Shelton,D.L., Tumas,D., Watanabe,C.K. and Wood,W.I.
Compositions and methods for the treatment of immune related
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|GGGAGCTGTCCAGAGCGCTGTGCCCCTGTCTGGTCTGGAGG
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A death-domain-containing receptor that mediates apoptosis Nature 384 (6607), 372-375 (1996)
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/gene="ws1-1"
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/db_xref="G1:1669691"
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Centre, Gunnels Wood Road,
Location/Qualifiers
                                                                                                                                                                            /db_xref="SWISS-PROT:093038"
/translation="MEGRPRGCAAVAAALLIVILIGARAQGGTRSPRCDCAGDEHKKIG
LFCCRGCPAGHYLKAPCTEPCGNSTCLVCPQDTFLAWENHHNSECARCQACDEQASQV
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/tissue_type="lymphoid"
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/db_xref="taxon:9606"
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/gene="ws1-1"
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RS Yu.G.-L., Ni.J., Gentz,R.L. au.
2 Death domain containing receptors
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Chaudiely, Chaudiely, Molecular B. Submitted (03-JAN-1997) Molecular B. Submitted (1705 NE Pacific Street,
                              Homo sapiens
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
1 (bases 1 to 1528)
Chaudhary.P.M. and Hood,L.E.
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                                                                                                                                                 GlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThrAspCysGlyThr 140
                                                                                                                                                                                                                                                                                                                                                    GAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCCAGGCTGGTTTGTGGAG
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AVADTROGCREGWTVECQVSGCVSSSPEYCQPCLDCGALHRHTRLLCSRRDTDCGTCL
LGFYEHGDGCVSCPTSTLGSCPERCAAVCGWRQMFWVQVLLAGLIVVPLLLGARLTYTY
RHCWPHKPLVTADEAGMEALTPEPATHLSPLDSAHTLLAFPDSSEKICTVQLLCGNSWT
PGYPETQEALCPQVTWSWDQLPSRALGPAAAPTLSPESPAGSFAMMLOPGPQLYDVMD
AVPARRWKEFVETLGLREAEIEAVEVEIGRFRDQQYEMLKRWRQQQPAGLGAVYAALE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /gene="DDR3"
/gene="DDR3"
/function="mediates apoptosis when cross-linked"
/function="mediates apoptosis when cross-linked"
/note="A TNFR1-related death-domain containing receptor;
/note="A TNFR1-related death-domain conta
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/map="1p36"
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<1. .1238
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/db_xref="taxon:9606"
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                                                          CAGGGCGCACTCGTAGCCCCAGGTGTGACTGTGCCGGTGACTTCCACAAGAAGATTGGT
                                                                                GlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHisLysLysIleGly
LeuPheCysCysArgGlyCysProAlaGlyHisTyrLeuLysAlaProCysThrGluPro
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1 (bases 1 to 1557)

Bodmer, J. L., Burns, K., Schneider, P., Hofmann, K., Steiner, V., Thome, M., Bornand, T., Hahne, M., Schroter, M., Becker, K., Wilson, A., French, L. E., Browning, J. L., MacDonald, R. and Tschopp, J.

TRAMP, a novel apoptosis-mediating receptor with sequence homology to tumor necrosis factor receptor 1 and Fas(Apo-1/CD95)

Immunity 6 (1), 79-88 (1997)
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                                                                                                                                                                                                                                                                                                                                                                                 /product="apoptosis-mediating receptor TRAMP"
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TCLLGFYEHGDGCVSCPTSTLGSCPERCAAVCGWQNEWQVLLAGLIVPLLLGATLF
TYRRIGWPHKPLVTADEAGMEALTPPPATHLSPLDSAHTLLAPPDSSEKICTVQLVGN
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VMDAVPAARWKEEYVTILGLREAEIEAVEVEIGRFRDQQYEMLKRWRQQQPAGLGAVYA
ALERMGLDGCVEDLRSRLQRGP"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /function="mediates apoptosis"
/note="contains a death domain; similar to TNF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 function="activates NF-kB"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /tissue_type="fetal lung"
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/db_xref="taxon:9606"
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-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                    This cDNA clone codes for human death domain containing receptor DR3 (see AAW31517), a novel member of the tumour necrosis factor receptor family. It was isolated from a HUVEC cDNA library. The receptor family. It was isolated from a HUVEC cDNA (see AAR89426) was isolated from a human testis tumour cDNA library. The genes consisted in cDNA libraries of foetal liver, foetal brain, tonsil and leukocyte. Nucleic acids encoding full-length or mature DR3, or the extracellular, transmembrane, intraclelular or especially the death domain of DR3, can be used to produce recombinant polypeptides in transformed host cells. These polypeptides can be used to treat diseases and disorders associated with increased apoptosis and for treating inflammatory diseases and disorders.
                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1254 BP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Death domain containing receptor polypeptide(s) DR3 and DR3-V1 - for activation of apoptosis and NF-kappaB, antagonists can be used
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                                     GluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGlyTrpPheValGlu
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                                                                                    HisAsnSerGluCysAlaArgCysGlnAlaCysAspGluGlnAlaSerGlnValAlaLeu
                                                                                                                                   CysGlyAsnSerThrCysLeuValCysProGlnAspThrPheLeuAlaTrpGluAsnHis
                                                                       CATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGCCTCCCAGGTGGCGCTG
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12-MAR-1996;
17-OCT-1996;
                                                         The invention provides nucleotide sequences encoding death domain containing receptor polypeptides DR3-V1 and DR3. The DR3-V1 cDNA clone is contained in ATCC deposition No. 97456 and the DR3 cDNA clone is contained in ATCC deposition No. 9757. Recombinant vectors comprising the nucleic acid sequences and optionally the leader sequences are used for the recombinant production of the proteins. The present sequence represents a cDNA encoding the death domain containing
    Sequence
                                                  receptor
                                                                                                                                                                                                                                                               Claim 6;
                                                                                                                                                                                                                                                                                                 New death domain containing receptor and recombinant vector optionally comprising leader sequence
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P-PSDB; AAW95538.
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(UNMI) UNIV MICHIGAN.
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96US-0028711.
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201 A; 420
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Pred. No.: Score: Alignment Scores:

7.96e-72 1038.00

Length: Matches:

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(YUGG/)
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                                                                                                                22-APR-1999;
28-MAY-1999;
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) HUMAN GENOME SCI INC.
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) GENTZ R L.
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Treating graft-versus-host disease, cancer, immunodeficiency or autoimmune disease comprising administering an antibody to Death Containing Receptor proteins and a second therapeutic agent
                                                                                             Example 2B; Fig
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DB; AAB36265.
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                                                                                           273pp; English.
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                                                                                                                                                                   Dillon PJ,
                                                                                                                                                                  Dixit VM;
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an h Domain

The present invention provides the protein and coding sequences for two death domain containing receptors, designated DR3 and DR3-V1. These receptors are involved in apoptosis, and the sequences given can be used in the treatment of cancers, infections, cardiovascular disorders such as arrhythmias, ischaemia, aneurysms, arterial occlusive diseases, embolisms and congenital heart defects, neurodegenerative diseases including Alzheimer's and Parkinson's diseases, autoimmune disease such as multiple solerosis, arthritis, diabetes, Graves' disease, asthma and psoriasis, to promote angiogenesis and wound healing.

Sequence 1254 B₽; 201 A; 420 Ç 407 G; 226 7 0 other;

Alignment Pred. No.:

No.:

Score:

Best Local S Query Match:

Percent Similarity: Best Local Similarity:

7.96e-72 1038.00 100.00% 100.00% 100.00%

Length: Matches: Conservative:

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US-09-993-234-6\_COPY\_25\_198 (1-174)

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Pred. No.:
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Query Match:
                                                                                                     Percent Similarity:
                                                                                                                                                                                                                                cDNA clone FH20.57 (ATCC 55820) codes for a novel polypeptide (see w26709), designated Apo-3, that stimulates or induces apoptotic activity in mammalian cells. It was isolated from a human foetal heart cDNA library by screening with probes (see T91183-84) based on an EST sequence (GenBank locus W7194) that showed homology to the intracellular domain of human TNFR1 and CD95. Amino acid residues 1-181 of Apo-3 are identical to another novel apoptosis polypeptide, Apo-2LI (see W26708). Nucleic acids encoding Apo-3 can be used diagnostically for tissue-specific typing and to produce recombinant Apo-3 polypeptides, especially the extracellular domain (amino acids 1-198) or death domain (amino acids 1-199) or death domain (amino acids 1-190) or d
                                                                                                                                                                                                 Sequence 1634 BP;
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     Diagnosing rheumatoid arthritis by probing digested human genomic or comparing expression of mrNA or polypeptide of a region of transmembrane protein \cdot
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P-PSDB;
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                                                                                  (SHIO/) SHIOZAWA
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                     Homo
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27-MAR-2001;
30-MAR-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Genomes, particularly DR3 genomic DNA, participating arthritis via mutation, useful in evaluating disease possibility and providing therapy and remedies -
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                                                                                                                                                                                                                                                                                                                                                              Sequence
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DB; AAO17879.
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18-FEB-2000;
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21-MAR-2000;
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20-JUL-1999
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15-SEP-1999
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01-DEC-1999
01-DEC-1999
01-DEC-1999
05-JAN-20000
11-FEB-2000
             Thirty three nucleic acids encoding PRO polypeptides which are useful in the diagnosis and treatment of immune related disorders, e.g. systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis,
                                                         P-PSDB;
                                                                                      Wood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; PRO; antiinflammatory; dermatological; antiarthritic; antirheumatic; cardiant; antianaemic; immunosuppressive; antithyrantidiabetic; noctropic; neuroprotective; hepatotropic; virucide; antiallergic; antiasthmatic; immune related disorder; hepatobiliary disease; autoimmune disease; allergy; ss.
                                                                                                                            (GETH )
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DB; AAB50918.
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                                                                                               NJ, Baker KP, Chan
Henzel W, Kabakoff
                                                                                                                                              99WO-US30911
2000WO-US00376
2000WO-US003565
2000WO-US03565
2000WO-US04341
2000WO-US04341
2000WO-US04414
2000WO-US04414
2000WO-US04884
2000WO-US06884
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99US-0146222
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                                                                                            Goddard A,
Shelton D
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                                                                                                      Godowski PJ,
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                                                                                            Tumas D,
                                                                                           , Gurney
Watanabe
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Alignment Scores: Pred. No.:
                                                                                                                                                                                                                            immune-mediated renal disease, demyelinating diseases of the central care immune-mediated renal disease, demyelinating diseases of the central care polyneuropathy or Guillain-Barre syndrome, and chronic demyelinating polyneuropathy or Guillain-Barre syndrome, and chronic inflammatory demyelinating polyneuropathy), hepatobiliary diseases:

( such as infectious, autoimmune chronic active hepatitis, primary billammatory bowel disease, gluten-sensitive enteropathy and Whipple's disease, autoimmune or immune-mediated skin diseases (such as bullous skin diseases, erythema multiforme, contact dermatitis, psoriasis), allergic diseases such as asthma, allergic rhinitis, atopic dermatitis, food hypersensitivity and urticaria), immunological diseases of the lung (such as eosinophilic pneumonias, idiopathic pulmonary fibrosis and hypersensitivity pneumonitis), transplantation associated diseases including graft rejection and graft-versus-host diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence is one of thirty three nucleic acids encoding PRO polypeptides. The PRO polypeptides, anti-PRO antibodies, agonists and antagonists are useful for treating and diagnosing immune related disorders such as systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis, juvenile chronic arthritis, spondyloarthropathies, systemic sclerosis, idiopathic inflammatory myopathies, Sjogren's syndrome, systemic vasculitis, sarcoidosis, autoimmune haemolytic anaemia, autoimmune thrombocytopaenia, thyroiditis, diabetes mellitus, immune-mediated renal disease, demyelinating diseases of the central anaemia control of the control of t
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US-09-993-234-6\_COPY\_25\_198 (1-174) x AAC91477 (1-1662) DB: Query Percent Similarity: Best Local Similarity: Match: 1.1e-71 1038.00 100.00% 100.00% 100.00% 22 Length: Matches: Conservative: Gaps: Mismatches: Indels: 1662 174 0 0 0

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294

355 61 CATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGCCTCCCAGGTGGCGCTG HisasnSerGluCysalaargCysGlnalaCysaspGluGlnalaSerGlnValalaLeu

81 GluAsnCysSerAlaValAlaAspThrArgCysGlyCysLySProGlyTrpPheValGlu 100

475 101 415 

Ωy 밁 Š 밁 Ω 밁 Ş 밁

В

밁 δÃ 535 121 GlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThrAspCysGlyThr 140 594

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δÃ 밁 655 161 GGGAGCTGTCCAGAGCGCTGTGCCGCTGTCTGTGGCTGGAGG GlySerCysProGluArgCysAlaAlaValCysGlyTrpArg 174 696

and

diabetes mellitus

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RESULT 8
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                      Score:
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                                                             Alignment
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11 MAR-1999
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22 JUN-1999
22 JUL-1999
26 JUL-1999
26 JUL-1999
27 AUG-1999
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Marsters SA,
Watanabe CK,
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01-DEC-1999;
05-JAN-2000;
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01-SEP-1999;
15-SEP-1999;
                                                                                                                                    The present invention relates to the isolation of novel human PRO polypeptides and the polynucleotide sequences encoding them. The PRO polypeptides, agonists, antagonists or anti-PRO antibodies are useful for treating benign or malignant tumours (e.g. renal, kidney, bladder, breast, etc), leukaemias and lymphoid malignancies, other disorders such as neuronal, glial, astrocytal, hypothalamic, glandular, macrophagal, stromal and blastocoelic disorders, inflammatory, immune and angiogenic disorders. The polynucleotide sequences are also useful in gene therapy. hMR40254-ABR40288 encode for the human PRO polypeptides of the invention.
                                                                                                                                                                                                                                                                                                                                                                                          Thirty five nucleic acids encoding PRO polypeptides, useful for treating benign or malignant tumours, leukaemias and lymphoid malignancies, inflammatory, angiogenic and immunologic disorders
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; PRO; benign tumour; malignant tumour; lymphoid malignancy; leukaemia; neuronal disorder; stromal disorder; blastocoelic disorinflammatory disorder; immune disorder; anglogenic disorder; gene therapy; cytostatic; neuroprotective; gene; ss.
                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                            Claim
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      Similarity:
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DB; AAU86139.
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99WO-US20111.
99WO-US21090.
99WO-US28313.
99WO-US28313.
99WO-US28634.
2000WO-US00219.
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Pan J, Pitti Wood WI;
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i RM, Roy MA,
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Query Match:
05-JUN-2000;
05-JUN-2000;
18-SEP-2000;
18-SEP-2000;
20-SEP-2000;
20-SEP-2000;
20-SEP-2000;
20-SEP-2000;
22-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                    stomach; lung; prostate; pancreas; carcinoma; antitumour; cancerous;
cytostatic; gene therapy; antineoplastic; Wilm's tumour; adenocarcinoma;
                                                                                                                                                                                                                                                                                                                                                                            stomach;
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                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                     gene;
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stomach; lung;
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2000US-209473P.

2000US-209531P.

2000US-233133P.

2000US-233617P.

2000US-234009P.

2000US-234052P.

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CC The present invention describes a method (M1) for screening for an CC anti-neoplastic agent. The method involves exposing cells to a chemical CC agent to be tested for anti-neoplastic activity, determining a change in CC expression of at least one gene (I) of a signature gene set, where (I) CC comprises a sequence (S) selected from 8447 sequences (given in ABL61664 CC to ABL70110), or is at least 95% identical to (S), where a change in CC expression is indicative of anti-neoplastic activity. (I) has cytostatic CC activity and can be used in gene therapy. MI can be used for screening CC an anti-neoplastic agent, and can be used for producting a product which CC is the data collected with respect to the anti-neoplastic agent as a CC result of MI, and the data is sufficient to convey the chemical CC structure and/or properties of the agent. MI can be used in the CC treatment of cancer such as colon, breast, stomach, lung, thyroid, CC adenocarcinoma, carcinoma, clear cell cancer, infiltrating ductal cancer, infiltrating ductal cancer, squamous cell carcinoma, neuroendocrine CC carcinoma, papillary carcinoma and Wilm's tumour.
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      Sequence 1743
                                                                                                                                                                                                                                                                                                                      Screening for anti-neoplastic agent involves exposing cells to chemical agent to be tested for anti-neoplastic activity, and determining a change in expression of a gene of a signature generation.
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Score: Pred. No.: Alignment

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Length:
Matches:
Conservative:
Mismatches:

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             12-MAR-1996;
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                                                                                                       Key
                                                                            mat_peptide
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                                         18-SEP-1997.
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                                                                                                                                               Death domain
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GENOME SCI
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                           96WO-US16849
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This cDNA clone, deposited as ATCC 97456, codes for human death domain containing receptor DR3-V1 (see AAW31516), a novel member of the tumour necrosis factor receptor family. It was isolated from a cDNA library derived from cells of a human testis tumour. Related death domain containing receptor DR3 cDNA (see AAT89427) was isolated from a HUVEC cDNA library. The genes have also been identified in cDNA libraries of foetal liver, foetal brain, tonsil and leukocyte. Nucleic acids encoding full-length or mature DR3-V1 can be used to produce recombinant polypeptides in transformed host cells. These polypeptides can be used to treat diseases and disorders associated with increased apoptosis. Antagonists, such as antibodies raised against DR3-V1, can be used to treat diseases and disorders associated with increased apoptosis and for treating inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 2; Page 71-73; 108pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Death domain containing receptor polypeptide(s) DR3 and DR3-V1 - for activation of apoptosis and NF-kappaB, antagonists can be used to treat inflammatory diseases
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GlySerCysProGluArgCysAlaAlaValCysGlyTrpArg
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                                                                                                                                           CysGlnValSerGlnCysValSerSerSerProPheTyrCysGlnProCysLeuAspCys
                          TGCCTGCCTGGCTTCTATGAACATGGCGATGGCTGCGTGTCCTGCCCCACGAGCACCCTG
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                                                                                                                                                                                                                                                                                                                            06-FEB-1997;
12-MAR-1996;
17-OCT-1996;
                                                                                                                                                   The invention provides nucleotide sequences encoding death domain containing receptor polypeptides DR3-V1 and DR3. The DR3-V1 cDNA clone is contained in ATCC deposition No. 97456 and the DR3 cDNA clone is contained in ATCC deposition No. 97757, Recombinant vectors comprising the nucleic acid sequences and optionally the leader sequences are used for the recombinant production of the proteins. The present sequence represents a cDNA encoding the death domain containing
                                                                                                                                                                                                                                                                                 WPI;
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                                                                                                                           Sequence
                                                                                                                                                                                                                            Claim
                                                                                                                                                                                                                                           New death domain containing receptor and recombinant vector optionally comprising leader sequence
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96US-0013285.
96US-0028711.
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        Treating graft-versus-host disease, cancer, immunodeficiency or autoimmune disease comprising administering an antibody to Death Containing Receptor proteins and a second therapeutic agent -
                                                                                              P-PSDB;
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autoimmune disorder;
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Apoptosis inducing receptor; AIR protein; Type I transmembrane protein; tumour cell

protein;

cell

human; cell death regulator; death; autoimmune disease;

Human apoptosis inducing receptor coding sequence

20-AUG-1998 AAV28700;

entry)

AAV28700 standard;

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                                                                                                                                                                                                                                                                                          This sequence encodes the human apoptosis inducing receptor (AIR) of the invention. AIR is a Type I transmembrane protein, soluble forms of which can be used to regulate cell death in a therapeutic setting. Soluble AIR can also be used in vitro to block apoptosis or AIR-expressing cells, or to screen agonists or antagonists of AIR activity. The cytoplasmic domain of AIR can be used to develop assays for inhibitors of AIR-induced cell death, which is useful to regulate cell death in a therapeutic setting as well as in vitro. Agonists of AIR activity can be used to kill tumour cells that express AIR, or T cells expressing AIR in autoimmune diseases.
                                                                                                                                                                                                                                                                                                                                                                           Claim
                                                                                                                                                                                                                                                                                                                                                                                         DNA encoding apoptosis inducing receptor - which is transmembrane protein, useful for regulating cell de
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RESULT 14
AAF83770
                                  Composition for treatment of T-cell mediated disease e.g. an cancer comprises a biologically active TR3-specific binding especially a monoclonal antibody -
                                                                                                    WPI; 2001-343711/36
P-PSDB; AAB84941.
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 Disclosure; Page 72;
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/note= "this codon has an apparent 1 nucleotide
deletion which alters the reading frame"
/transl_except= "(pos:558..559, aa:Cys)"
/note= "this codon has an apparent 1 nucleotide
deletion which alters the reading frame"
/transl_except= "(pos:538.539, aa:Leu)"
/note= "this codon has an apparent 1 nucleotide
deletion which alters the reading frame"
/transl_except= "(pos:718.719, aa:Met)"
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The invention relates to a composition comprising a biologically active TR3-specific binding agent (I) that binds to TR3 and inhibits the

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RESULT 15
ABL67258
ID ABL6
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AC ABL6
XX
AC ABL6
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Thyr
DT 15-M
XX
XX
Huma
KW Huma
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Best Local Similarity:
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               Human; cancer;
                                                                  15-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              proliferation of cells expressing TR3. (I) identified by the methods are useful for treating a subject suspected of having a disease associated with a proliferation of cells expressing TR3 especially leukemias or lymphomas or a T-cell mediated disease especially autoimmune diseases such as myasthenia gravis, systemic lupus erythamatosus, rheumatoid arthritis, diabetes, multiple sclerosis, sarcoidosis, myocarditis, thyroiditis and tumours. (I) is also useful for treating a subject suspected of having graft-versus-host disease, rejection of a transplanted organ such as heart, liver, lung, kidney, pancreas, bowel, skin or an appendage, or inflammatory diseases, allergies and contact dermatitis. The present sequence represents the nucleotide sequence of having a cone
                                                                                           ABL67258
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||GGAGCGT-CCAGAGCGCTGTGCCCGCTGTCTGTGGCTGGAGG
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colon; breast; ovary; oesophagus; kidney;
prostate; pancreas; carcinoma; antitumour
                                     related gene sequence
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27-SEP-2000;
28-SEP-2000;
28-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                                                                               cytostatic;
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2000US-233617P.
2000US-234009P.
                                                                                                                                                                                                                                                                                                                                                                                              gene therapy; antineoplastic;
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2000US-234923P
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#### AVALON PHARM.

Augustus Weaver 2 įχ Carter č, Ebner R, Endress ç, Horrigan S

# 2002-188264/24.

Claim

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44pp;

English

chemical determining Screening

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for anti-neoplastic agent involves exposing cells agent to be tested for anti-neoplastic activity, ar a change in expression of a gene of a signature

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set

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The present invention describes a method (M1) for screening for an anti-neoplastic agent. The method involves exposing cells to a chemical agent to be tested for anti-neoplastic activity, determining a change is expression of at least one gene (I) of a signature gene set, where (I) comprises a sequence (S) selected from 8447 sequences (given in ABL6166. expression of at least comprises a sequence (s to ABL70110), or is at change ABL61664 'n

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   expression is indicative of anti-neoplastic activity. (I) has cytostatic activity and can be used in gene therapy. M1 can be used for screening an anti-neoplastic agent, and can be used for producing a product which is the data collected with respect to the anti-neoplastic agent as a result of M1, and the data is sufficient to convey the chemical structure and/or properties of the agent. M1 can be used in the treatment of cancer such as colon, breast, stomach, lung, thyroid, oesophageal, ovarian, kidney, prostate or pancreatic cancer, adenocarcinoma, carcinoma, clear cell cancer, infiltrating ductal cancer, squamous cell carcinoma, neuroendocrine
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                                                                   GlySerCysProGluArgCysAlaAlaValCysGlyTrpArg 174
                                                                                                                                                                                         TGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGCCAACCATGCCTAGACTGC
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OM protein - nucleic search, using frame\_plus\_p2n model

GenCore version 5.1.4\_p5\_4578 Copyright (c) 1993 - 2003 Compugen Ltd.

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Result
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-LOOPCL-0 -LOOPEXT-0 -UNITS-bits -START-1 -END--1 -MATRIX-blosum62
-TRANS-human40.cdi -LIST-45 -DOCALIGN-200 -THR_SCORE-pct -THR_MAX=1.00
-THR_MIN-0 -ALIGN-15 -MODE-LOCAL -OUTFMT-pto -NORM-ext -HEAPSIZE-500 -MINLEN-0
-MAXLEN-200000000 -USER-US09993234_@CGN_11_249_@runat_27032003_115459_15466
-NCPU-6 -ICPU-3 -NO_XLPXY -NO_MMAP -LARGEQUERY -NUS_SCORES-0 -WAIT -LONGLOG
-DEV_TIMEOUT-120 -WARN_TIMEOUT-30 -THREADS-1 -XGAPOP-10 -XGAPEXT-0.5 -FGAPOP-6
-FGAPEXT-7 -YGAPOP-10 -YGAPEXT-0.5 -DELOP-6 -DELEXT-7
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Maximum Match 100%
Listing first 45 summaries
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-MODEL=frame+_p2n.model -DEV=xlh
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Fgapop 6.0 , F
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    /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq: *
    /cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq: *
    /cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq: *
    /cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq: *
    /cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq: *
    /cgn2_6/ptodata/1/pubpna/US09_PUBCOMB.seq: *
    /cgn2_6/ptodata/1/pubpna/US09_PUBCOMB.seq: *
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/cgn2_6/ptodata/1/pubpna/PCT_NEW_PUB.seq:*
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US-10-081-280-9
US-10-112-793-9
US-10-112-193-11
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Sequence 3, Appli
Sequence 9, Appli
Sequence 9, Appli
Sequence 11, Appl
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## ALIGNMENTS

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US-09-33-966-3
; Sequence 3, Application US/09333966
; Patent No. US20020009773A1
; GENERAL INFORMATION:
APPLICANT: Yu, Guo-Liang
APPLICANT: Ni, Jian
APPLICANT: Dixit, Vishva
APPLICANT: Dixit, Vishva
APPLICANT: Dillon, Patrick J.
TITLE OF INVENTION: Death Domain Containing Receptors
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSE: Sterne, Kessler, Goldstein & Fox, P.L.L.C.
STREET: 1100 New York Ave., NW, Suite 600
CITY: Washington
STATE: DC
COUVERY: USA
ZIP: 20005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NOMBER: US/09/333,966
FILING DATE:
CLASSIFICATION:
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PRIOR APPLICATION DATA:

US/08/815,469

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US-09-993-234-6_COPY_25_198 (1-174) x US-09-333-966-3 (1-1254)
 161 GlySerCysProGluArgCysAlaAlaValCysGlyTrpArg 174
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                                                                                                                                                     CysGlnValSerGlnCysValSerSerSerProPheTyrCysGlnProCysLeuAspCys 120
                         TGCCTGCCTGGCTTCTATGAACATGGCGATGGCTGCGTGTCCTGCCCCCACGAGCACCCTG
                                         GlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThrAspCysGlyThr 140
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Best Local Similarity:
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US-10-081-280-9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 415/952-9881
TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
281
                                                                                                                                                               221
                                                                                                                        21 LeuPheCysCysArgGlyCysProAlaGlyHisTyrLeuLysAlaProCysThrGluPro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/829,270
FILING DATE: 31-Mar-1997
ATTORNEY/AGENT INFORMATION:
NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007R1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                            CTGTTTTGTTGCAGAGGCTGCCCAGCGGGGCACTACCTGAAGGCCCCTTGCACGGAGCCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: 415/225-5416
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 1
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TITLE OF INVENTION: Apo-2 LI AND
NUMBER OF SEQUENCES: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
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TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/10/081,280 FILING DATE: 21-Feb-2002 CLASSIFICATION: <Unknown>
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CITY: South San Francisco
STATE: California
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Match:

Percent Similarity: Best Local Similarity:

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Score

Alignment Scores: US-09-333-966-3

No.:

MOLECULE TYPE: FEATURE:

TOPOLOGY:

both CDNA

NAME/KEY: LOCATION:

CDS

INFORMATION FOR SEQ ID NO: 3: SEQUENCE CHARACTERISTICS:

LENGTH: 1254 base pairs
TYPE: nucleic acid
STRANDEDNESS: double

TELECOMMUNICATION INFORMATION: TELEPHONE: 202-371-2600

NAME: Steffe, Eric K.
REGISTRATION NUMBER: 36
REFERENCE/DOCKET NUMBER:

36,688

APPLICATION NUMBER: US 60/013,285
FILING DATE: 12-MAR-1996
ATTORNEY/AGENT INFORMATION:

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/028,711
FILING DATE: 17-001-1996

FILING DATE: APPLICATION NUMBER: APPLICATION NUMBER:

PRIOR APPLICATION DATA:

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253

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Percent Similarity:
Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                    TELEFAX: 650/952-9881 INFORMATION FOR SEQ ID NO: 9:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
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                                                                                                                                                                                                                                             APPLICATION NUMBER: US/10/112,793
FILING DATE: 28 Mar-2002
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/828,683A
FILING DATE: 31-Mar-1997
APPLICATION NUMBER: 08/625328
FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
ATTORNEY/AGENT INFORMATION:
NAME: MATSCHANG, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: 91007P1
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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MEDIUM TYPE: 3.5 inch, 1.44 Mb i
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
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TITLE OF INVENTION: Apo-2 LI AND Apo-3 POLYPEPTIDES
NUMBER OF SEQUENCES: 28
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                                                                                                                                         TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: South San Francisco
STATE: California
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US-10-112-193-11
; Sequence 11, Application US/101
; Publication No. US20030004313A1
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                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/10/112,193
FILING DATE: 28-Mar-2002
CLASSIFICATION: CURROWN>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            341
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            ATTORNEY/AGENT INFORMATION:
NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 i
                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Apo-3 POLYPEPTIDE NUMBER OF SEQUENCES: 15
                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Ashkenazi, Avi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GlySerCysProGluArgCysAlaAlaValCysGlyTrpArg 174
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FILING DATE: 11-Sep-1997
APPLICATION NUMBER: 60/026943
FILING DATE: 23-Sep-1996
                                                                                                                                                                                                                                                                              ZIP:
                                                                                                                                                                                                                                                                                          CITY: South San F.
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                          STREET:
REFERENCE/DOCKET NUMBER: P1052R1
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RESULT 5
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Best Local Similarity:
                                                                                                                                                    Sequence
Patent No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Alignment Scores: Pred. No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; SEQUENCE DESCRIPTION: SEQ US-10-112-193-11
                                                                                   equence 9, Application US/09993234
atent No. US20020146768A1
GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-2 LI AND Apo-3
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
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                                 STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STATE: California
                            COUNTRY: USA
                                                                           ADDRESSEE:
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TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
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Conservative:
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Best Local Similarity:
Query Match:
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161 GlySerCysProGluArgCysAlaAlaValCysGlyTrpArg
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                                               CysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysProThrSerThrLeu 160
                                                                                                                                 TGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGCCAACCATGCCTAGACTGC
                                                                                                                                               CysGlnValSerGlnCysValSerSerSerProPheTyrCysGlnProCysLeuAspCys
                                                                                                                                                                                 GAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCCAGGCTGGTTTGTGGAG
                                                                                                                                                                                             GluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGlyTrpPheValGlu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE DESCRIPTION: SEQ
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                                                                                                                                                                                                                                                                                                                                           LeuPheCysCysArgGlyCysProAlaGlyHisTyrLeuLysAlaProCysThrGluPro
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STRANDEDNESS: Single
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US-09-993-234-6_COPY_25_198 (1-174) x US-09-993-234-9 (1-1634)
                                                                                                                                                                                                                                                                                                                                         TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 1634 base pairs
1 GlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHisLysLysIleGly
                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007P1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/993,234
FILING DATE: 19-NO. US20020146768A1-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 08/828,683
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Marschang, Diane L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech)
                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 415/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: 415/225-5416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: <Unknown>
RAPPLICATION DATA:
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                                                                                                                                                                                                                                                                      NO:
                                                                                                               Conservative: Mismatches: Indels:
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Matches:
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US-09-993-234-6_COPY_25_198 (1-174) x US-09-954-531-1389 (1-1743)
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Query Match:
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US-09-954-531-1389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US/60/234,009
PRIOR FILING DATE: 2000-09-20
PRIOR APPLICATION NUMBER: US/60/234,034
PRIOR FILING DATE: 2000-09-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US/60/234,567
PRIOR FILING DATE: 2000-09-22
NUMBER OF SEQ ID NOS: 1392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/954,531
CURRENT FILING DATE: 2002-05-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Process for Identifying Anti-Cancer Therapeutic Agents Using Canc
TITLE OF INVENTION: Gene Sets
FILE REFERENCE: 689290-77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US/60/233,133
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
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                                                                                                                                                                                                                                                       GluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGlyTrpPheValGlu 100
                                                                                                                                                                                                                                                                                                          CysGlyAsnSerThrCysLeuValCysProGlnAspThrPheLeuAlaTrpGluAsnHis 60
                                                                                                                                                                 TGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGCCAACCATGCCTAGACTGC 500
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Best Local Similarity:
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                                   US-09-993-234-6_COPY_25_198 (1-174) x US-09-333-966-1 (1-1783)
                                                                                        Query Match:
                                                                                                                                                Score:
                                                                                                                                                             Alignment Scores: Pred. No.:
                                                                                                                                                                                                                     US-09-333-966-1
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US-09-333-966-1
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APPLICANT:
APPLICANT:
APPLICANT:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/028,711
FILING DATE: 17-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: ZUUUJ COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
TYPE: Floppy disk
                                                                                                                                                                                                                                                                        MOLECULE TYPE: FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 12-MAR-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: 202-371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Dixit, Vishva
APPLICANT: Gentz, Reiner L.
APPLICANT: Dillon, Patrick J.
TITLE OF INVENTION: Death Domain Containing Receptors
                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                      NAME/KEY:
LOCATION:
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                                                                                                                                                                                                                                                                                                              TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER:
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Matches:
Conservative:
Mismatches:
Indels:
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Percent Similarity:
Best Local Similari
Query Match:
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US-09-964-824A-292
                                                                                                                                                                                                                                           ; ORGANISM: Homo sapiens
US-09-964-824A-292
                                                                                                                                                                                                       Alignment Scores:
                                                                                                                                                                                                                                                                                                              SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence 292, Application US/09964824A
Patent No. US20020102531A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signatu
TITLE OF INVENTION: Sets
FILE REFERENCE: 689290-73
CURRENT APPLICATION NUMBER: US/09/964,824A
CURRENT FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: US/60/236,033
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR FILING DATE: 2000-09-28
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 GlySerCysProGluArgCysAlaAlaValCysGlyTrpArg 174
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                                                                                                                                    Similarity:
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93.68%
93.68%
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Conservative:
Mismatches:
Indels:
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US-10-081-280-5
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        APPLICATION NUMBER: US/08/829,270
APPLICATION NUMBER: US/08/829,270
FILLING DATE: 31-Mar-1997
ATTORNEY/AGENT INFORMATION:
NAME: MATSCHANG, DIANE L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007R1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-5416
TELEPHONE: 415/225-5416
TELEPAX: 415/952-9881
TELEPAX: 415/952-9881
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1438 base pairs
TYPE: NUCleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 5, Application US/10081280 Patent No. US20020165157Al GENERAL INFORMATION:
                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

COMPUTER: 3.5 inch, 1.44 Mb flopp
COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/10/081,280
FILING DATE: 21-Feb-2002
CLASSIFICATION: CUnknown>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Apo-2
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CITY: South San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
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Score: Pred.

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TOPOLOGY: Linear;
SEQUENCE DESCRIPTION:
US-10-081-280-5
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Best Local Similarity:
Query Match:
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Publication No. US20020192729A1
GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-2 LI AND
NUMBER OF SEQUENCES: 28
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                             ZIP: 94080

COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb 1

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: WinPatin (Genentech)
                       CURRENT APPLICATION DATA:
                                                                                                                                            CORRESPONDENCE ADDRESS:
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                                                                                                   ADDRESSEE: Genentech, Inc
STREET: 1 DNA Way
CITY: South San Francisco
STATE: California
   APPLICATION NUMBER: US/10/112,793 FILING DATE: 28-Mar-2002
                                                                                           COUNTRY: USA
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Percent Similarity:
Best Local Similarity:
Query Match:
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TELEPHONE: 650/225-5416
TELEPAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/828,683A
FILING DATE: 31-Mar-1997
APPLICATION NUMBER: 08/625328
FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
ATTORNEY/AGENT INFORMATION:
NAME: MAISCHARY, Diane L.
REGISTRATION NUMBER: 35,600
PREGEDENCE COCKET NUMBER: 35,600
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                                       GlySerCysProGluArgCysAlaAlaValCysGlyTrpArg
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TYPE: Nucleic Acid
STRANDEDNESS: Single
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
101 CysGlnValSerGlnCysValSerSerSerProPheTyrCysGlnProCysLeuAspCys 120
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                                  GAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCCCAGGCTGGTTTGTGGAG
                                                   GluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGlyTrpPheValGlu 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
                                                                                                  CATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGCCTCCCAGGTGGCGCTG
                                                                                                                     HisasnSerGluCysAlaArgCysGlnAlaCysAspGluGlnAlaSerGlnValAlaLeu
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
                                                                                                                                                                  TGCGGCAACTCCACCTGCCTTGTGTGTCCCCAAGACACCTTCTTGGCCTGGGAGAACCAC
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CURRENT APPLICATION DATA:
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MEDIUM TYPE: 3.5 1:
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ADDRESSEE: Genentech, Inc.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 1438 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 08/928,069
FILING DATE: 11-Sep-1997
APPLICATION NUMBER: 60/026943
FILING DATE: 23-Sep-1996
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FILING DATE: 28-Mar-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: 3.5 inch, 1.44 Mb 1
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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ZIP: 94080
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Matches:
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                                                        US-09-993-234-6_COPY_25_198 (1-174) x US-09-884-733-5 (1-1438)
                                                                                                        Query Match:
                                                                                                                   Percent Similarity:
Best Local Similarity:
                                                                                                                                                   Score:
                                                                                                                                                                                                       ; SEQUENCE DESCRIPTION: SEQ ID US-09-884-733-5
                                                                                                                                                               Pred. No.:
                                                                                                                                                                            Alignment Scores:
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US-09-884-733-5
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Patent No. US20020123116A1
                                                                                                                                                                                                                                                                                                      TELEFAX: 415/952-9881
TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 5:
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161 GlySerCysProGluArgCysAlaAlaValCysGlyTrpArg 174
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NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,6
                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-5416
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ADDRESSEE: Genentech,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Apo-2 Ligand Inhibitor NUMBER OF SEQUENCES: 9
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FILING DATE: 19-Jun-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: 15M PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
NT APPLICATION:
                                                                                                                                                                                                                                                            TYPE: Nucleic Acid
                                                                                                                                                                                                                                  TOPOLOGY: Linear
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                                                                                                                                                                                                                                             STRANDEDNESS: Single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: South San Francisco
                                                                                                                                                                                                                                                                          LENGTH: 1438 base pairs
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(BER: P1007
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Matches:
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RESULT 13
US-09-993-234-5
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Patent No. US20020146768A1
GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: App-2 LI AND App-3 POLYPEPTIDES
                                                              TELEFAX: 415/952-9881
TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 5:
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                                                                                                        FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: MAISCHANG, Diane L.
REGISTRATION NUMBER: 35,600
REGISTRATION NUMBER: P1007P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-5416
                                                                                                                                                                                                                                                                                               MEDIUM TYPE: 3.5 inch, 1.44 Mb
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch,
                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 460 Point San Bruno
CITY: South San Francisco
                               SEQUENCE CHARACTERISTICS:
LENGTH: 1438 base pairs
                                                                                                                                                                                                                                      PRIOR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GlySerCysProGluArgCysAlaAlaValCysGlyTrpArg 174
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                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/09/993,234
FILING DATE: 19-No. US20020146768A1-2001
CLASSIFICATION: <Unknown>
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                              STATE: California COUNTRY: USA
                                                                                                                                                                                                                    APPLICATION NUMBER: 08/828,683
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Pred. No.:
                                                                                 ; LENGTH: 10797
; TYPE: DNA
; ORGANISM: HOMO :
US-10-092-154-1577
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US-10-092-154-1577
JS-10-092-157, Application US/10092154
Publication No. US20030054375A1
GENERAL INFORMATION:
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  Best Local Similarity:
                Percent Similarity:
                            Score:
                                          Alignment Scores: Pred. No.:
                                                                                                                    CURRENT APPLICATION NUMBER: US/10/092,154
CURRENT FILING DATE: 2002-03-07
NUMBER OF SEQ ID NOS: 2003
Prior Application removed - See File Wrap)
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1577
LENGTH: 10797
                                                                                                                                                                                                                APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PC009C1
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Usery March:   71.10%	132	gHisThrArgLeuLeuCysSer	125	Qy
1.10%   Indels: 239	ଟ= r	nCysValSerSerSerProPheTyrCysGlnProCysLeuAspCysGlyAlaLe	16	Ωу
BETY METCH: 91.10% Indels: 429  1 GlnG1yGlyThrargSerFroArgCygAspCysAlaGlyAspCheHislyLySlegIy  2 GlnG1yGlyThrargSerFroArgCygAspCysAlaGlyAspCheHislyLySlegIy  2 LeuPhcysCysArgGlyCysProAla	CA 6	aValAlaAspThrArgCysGlyCysLysProGlyTrpPheValGluCysGlnVa.	<u> </u>	P 64
indels: 239  239-09-993-234-6_COPY_25_198 (1-174) x US-10-092-154-1577 (1-10797)  1 GlnGlyGlyThrArgSerProArgCysApgCysAlaGlyAspPheHislySlySileGly   GlnGlyGlyThrArgSerProArgCysArgGlySapCysAlaGlyAspPheHislySlySileGly   GlnGlyGlyThrArgSerProArgCysArgGlyGacTTGCCACAAGAAGATTGCT   GlnGlyGlyThrArgSerProArgCysArgGlyCysproAla	ი	ACTCCCCATCTCCCTGCACCCCCCCCCACCAGCC-TCCCAGGTGGCGCTGGAGAACTG	Ö	Db
indels: 239 is: 93124-6_COPY_25_198 (1-174) x US-10-092-154-1577 (1-10797)  1 GlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHislySLySileGly a Gaps: 239  21 LeupheCysCysArgGlyCysProAla	, o	CTTGGGATAATGCCTCTGCCTCTCC	7	Ov Db
LICY Match: 71.10% Gaps: 439  1.09-993-234-6_COPY_25_198 (1-174) x US-10-092-154-1577 (1-10797)  1. GlinglyGlyThrargSerProArgCysAspCysAlaClyAspPheHislySiySileGly and the control of the c	7		7	γ
HERY MATCH:  71.10%  1 Glaps: 29 29 21 GlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHislySlySileGly 21 GlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHislySlySileGly 22 LeuphecysCysArgGlyCysProAla	TCTCC 5	TCTCTTTTTAGGGTAGCCCTGTACCTGTCTGTCTT	92	뫄
Indels: 239	7		73	Q
Indels: 239	TCTG 5	TTCCTGGTGACCTTCCCATCTCCCATGTGCCTTGGCCTC	96	Db
Indels: 239	7		73	Qy
arry Match: 71.10% Indels: 239  3: 993-234-6_COPY_25_198 (1-174) x US-10-092-154-1577 (1-10797)  1 GlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHisLysLysIleGly 2  1 LeuPheCysCysArgGlyCysProAla	TTCTCT 5	CCACCCTTGCCAGAACCCTCCACCCTGATCCTCCTTCAGGGTGCCCTTGCCCC	80	В
LETY Match: 71.10% Gaps: 239  1 GlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHisLysLysIleGly 1 GlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHisLysLysIleGly 2   1   1   1   1   1   1   1   1   1	7		73	Qy
icry Match: 9.1.10% Gaps: 4  s-09-993-234-6_COPY_25_198 (1-174) x US-10-092-154-1577 (1-10797)  1 GlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHisLysLysIleGly	CCTGCA 5	AGGCCTTTCTGAAGGAAGTGGCTGGCTCGGGCCCAAACTTGGGGTGTGAGGGT	74	밁
lery Match: 9.1.10% Gaps: 4  1-09-993-234-6_COPY_25_198 (1-174) x US-10-092-154-1577 (1-10797)  1 GlnGlyGlyThrargSerProargCysAspCysAlaGlyAspPheHisLysLysIleGly	7		73	Qy
lidels: 239  1.10% Gaps: 4  1.109-993-234-6_COPY_25_198 (1-174) x US-10-092-154-1577 (1-10797)  1.11	CTAAGGA 5		89	DЬ
ery Match: 71.10% Indels: 239  1: 09-993-234-6_COPY_25_198 (1-174) x US-10-092-154-1577 (1-10797)  1 GlnGlyGlyThrargSerProargCysAspCysAlaGlyAspPheHisLysLysLleGly 2	7	GlnAlaCysAspGlu	68	Qy
ery Match: 71.10% Indels: 239 8: 9 Gaps: 4 1-09-993-234-6_COPY_25_198 (1-174) x US-10-092-154-1577 (1-10797) 1 GlnGlyGlyThrargSerProArgCysAspCysAlaGlyAspPheHisLysLysLeGly	υ σ	lCysProGlnAspThrPheLeuAlaTrpGluAsnHisHisAsnSerGluCysAl 	62	Оу
rery Match: 71.10% Indels: 239  8: 9 Gaps: 4  1-109-993-234-6_COPY_25_198 (1-174) x US-10-092-154-1577 (1-10797)  1 GlnGlyGlyThrargSerProArgCysAspCysAlaGlyAspPheHisLysLysLieGly 2	H-	ACAGGGCACTACCTGAAGGCCCCTTGCACGGAGCCCTGCGGC	56	В
ery Match: 71.10% Indels: 239  1 Gaps: 4  1 Gaps: 4  1 GlnGlyGlyThrargSerProArgCysAspCysAlaGlyAspPheHisLysLieGly   1   1   1   1   1   1   1   1   1	a 4	GlyHisTyrLeuLysAlaProCysThrGluProCysGly	30	Qy
ery Match: 71.10% Indels: 239  1. 109-993-234-6_COPY_25_198 (1-174) x US-10-092-154-1577 (1-10797)  1. GlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHisLysLieGly 2	CTGGTTC 5	aaagcaggtcggggttgctgggcagcccttctgcctgcct	50	Db
ery Match: 71.10% Indels: 239 ;: 91.10% Gaps: 4 ;:-09-993-234-6_COPY_25_198 (1-174) x US-10-092-154-1577 (1-10797)  1	2		29	Qy
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ery Match: 71.10% Indels: 239 : 9 Gaps: 4 -09-993-234-6_COPY_25_198 (1-174) x US-10-092-154-1577 (1-10797) 1 GlnGlyGlyThrArgSerPToArgCysAspCysAlaGlyAspPheHisLysLysLieGly 2	CATGGG 5	STTTTGTTGCAGAGGCTGCCCAGCGGGTAAGTGGCCACAGGGGTGGGAG	32	Db
ery Match: 71.10% Indels: 239 : Gaps: 4 -09-993-234-6_COPY_25_198 (1-174) x US-10-092-154-1577 (1-10797) 1 GlnGlyGlyThrargSerProArgCysAspCysAlaGlyAspPheHisLysLysIleGly 2	2	LeuPheCysCysArgGlyCysProAla	21	Qy
ery Match: 71.10% Indels: 239 : Gaps: 4 -09-993-234-6_COPY_25_198 (1-174) x US-10-092-154-1577 (1-10797) 1 GlnGlyGlyThrargSerProArgCysAspCysAlaGlyAspPheHisLysLysIleGly 2	TGGT 5	AGGCCGCACTCGTAGCCCCAGGTGTG	26	р
ery Match: 71.10% Indels: 239 ; Gaps: 4 -09-993-234-6_COPY_25_198 (1-174) x US-10-092-154-1577 (1-10797	eGly 2	GlnGlyGlyThrArgSerProArgCysA	1	Qy
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Qy 48 lCysProGlnAspThrPheLeuAlaTrpGluAsnHisHisAsnSerGluCysAlaArgCy 68 	Oy 30GlyHisTyrLeuLysAlaProCysThrGluProCysGlyAsnSerThrCysLeuVa 48	Db 5508 TGGAAAGCAGGTCGGGGGTTGCTGGGCAGCCCTCTGCCTGACCCCTGCCTG	Db 5448 GGTAGGGGTAGGCTGACAGAGAAGTAGGGGAGCTGGAGAGAAAGAGGGAAGGGAGGG	Db 5388 GCAGGCAGGGCTGGAGAGGTGGCGGGGAGGCCCGGGGAGGTAAGAGGAGGCTGGCAGGGGA 5447	ОУ 29 29	QY 21 LeuPheCysCysArgGlyCysProAla 29	Db 5268 CAGGGCGCACTCGTAGCCCCCAGGTGTGACTGTGCCGGTGACTTCCACAAGAAGATTGGT 5327	Qy 1 GlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHisLysLysIleGly 20	US-09-993-234-6_COPY_25_198 (1-174) x US-09-764-847-1577 (1-10797)	Alignment Scores:   1.29e-66   Length:   10797     Score:   738.00   Matches:   161     Percent Similarity:   39.95%   Conservative: 0     Best Local Similarity:   39.95%   Mismatches:   5     Query Match:   71.10%   Indels:   239     DB:   10   Gaps:   4	Prior ap NUMBER C SOFTWARE SEQ ID NG SEQ ID NG LENGTH: TYPE: D ORGANIS ORGANIS	US-09-764-847-1577 ; Sequence 1577, Application US/09764847 ; Patent No. US20020132767A1 ; GENERAL INFORMATION: ; APPLICANT: Rosen et al. ; TITLE OF INVENTION Nucleic Acids, Proteins, and Antibodies ; FILE REFERENCE: PC009 ; CURRENT APPLICATION NUMBER: US/09/764,847	DD 6442 ATGGAGG 6448 RESULT 15	172 yTrpArg 174	Qy 152 sValSerCysProThrSerThrLeuGlySerCysProGluArgCysAlaAlaValCysGl 172	Qy . 133 -ArgArgAspThrAspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCy 152	Db 6286 CTCCCTGCCTGACCCACTCCTGTCCCATGGTGACGCATGCCTCTCCTGGATTGCAGGTTC 6345	133

Q	68	sGlnAlaCysAspGlu 73
밁	5688	CCAGGCCTGTGATGAGCAGGGTGAGGGGCTTCTCAGTGCTTGGCAGGGAGTTCCTAAGGA 5747
Ωy	73	73
밁	5748	CAGGCCTTTCTGAAGGAAGTGGCTGGCTCGGGCCCAAACTTGGGGTGTGAGGGTCCTGCA 5807
QΥ	73	73 ,
B	5808	CCCACCCTTGCCAGAACCCTCCACCCTGATCCTCCTTCAGGGTGCCCTTGCCCCCTTCTCT 5867
δ	73	73
망	5868	CTTCCTGGTGACCTTCCCATCTCTCCATGTGCCTTGGCCTCTGGTCGGCCTTAATCTCTG 5927
Qy	73	73
ф	5928	AGCTTCTCTTTTTTAGGGTAGCCCTGTACCTGTCTGTCTTTCGCCTATTTCTGTCTCC 5987
δ	73	73
망	5988	ATTATCTTGGGATAATGCCTCTGCCTCTCCATGGGAGCCTTTGGCCCTGACTAACTCTCC 6047
δĀ	74	GlnAlaSerGlnValAlaLeuGluAsnCysSerAl 85
망	6048	ACTCCCCATCTCCCTGCACCCCCACCAGCC-TCCCAGGTGGCGCTGGAGAACTGTTCAGC 6106
Qγ	85	aValAlaAspThrArgCysGlyCysLysProGlyTrpPheValGluCysGlnValSerg1 105
В	6107	AGTGGCCGACACCCGCTGTGGCTGTAAGCCAGGCTGGTTTGTGGGAGTGCCAGGTCAGCCA 6166
γ	105	nCysValSerSerSerProPheTyrCysGlnProCysLeuAspCysGlyAlaLeuHisAr 125
B	6167	ATGTGTCAGCAGTTCACCCTTCTACTGCCAACCATGCCTAGACTGCGGGGCCCTGCACCG 6226
VΩ	125	gHisThrArgLeuLeuCysSer 132
문	6227	CCACACACGGCTACTCTGTGA-GTACCCCCACCCAGGGCTCTCTACTCCCAGACCCCCTT 6285
γ	132	132
망	6286	CTCCCTGCCTGACCCACTCCTGTCCCATGGTGACGCATGCCTCCTGCATTGCAGGTTC 6345
δ	133	-ArgArgAspThrAspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCy 152
ф	6346	CCGCAGAGATACTGACTGTGGGACCTGCCTGCCTGCCTTCTATGAACATGGCGATGGCTG 6405
δĀ	152	hrLeuGlySerCysProGluArgCysAlaAlaValCys
망	6406	CGTGTCCTGCCCCACGTAATTCCTA
δ	172	YTrpArg 174
Вb	6442	ATGGAGG 6448
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Command line parameters:

-MODEL-frame+_p2n.model -DEV-xlh
-Q-/cgn2_1/USPT0_spool/US09993234/runat_27032003_115456_15362/app_query.fasta_1.2346
-DB-EST -OFMT-fastap -SUFFIX=rst -MINMATCH=0.1 -LOOPCL=0 -LOOPEXT=0
-UNITS-bits -START=1 -END--1 -MATRIX-blosum62 -TRANS-human40.cdi -LIST=45
-DOCALIGN=200 -THR_SCORE-pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15 -MODE-LOCAL
-OUTFMT-pto -NORM-ext -HEAPSIZE=500 -MILLEN=0 -MAXLEN=2000000000
-USER-US0993334_eCGN_11_4749_@runat_27032003_115456_15362 -MCPU=6 -ICPU=3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120
-MARN_TIMEOUT=30 -THREADS=1 -XGAPD=10 -XGAPEXT=0.5 -FGAPDP=6 -FGAPEXT=7
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

### SUMMARIES

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$\alpha \cdot \cdot \mu \mu$	323 296 292 277.5 268.5 214 211	436.5 420.5 416 405 395.5 389.5 360.5	71 88 99 99 99 99 99 99 99 99 99 99 99 99	Score
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57787	2561449	775 322 330 316 639 393	767 6638 6638 5562 5562 5562 5562 5563 5563 5563 5563	Length
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AI1193 BF164 BI555 BI769 AV664		677 124 141 675 363 851	B19994 AW26861 A1424936 A1807913 A1140043 BM7807913 BM7817430 BM75437 BM71397 BM71397 BM7103624 BM77079 BM710769 BM710769 BM71079 BM71	ID
19338 uf03d04.y 164835 60177775 555040 60323655 769006 60305816 664984 AV664984	6378 yolle 6424 yollel 9739 yn60h0 692368 UI-E 872434 BB87 695862 UI-E	67742 6027868 1249 op73c01. 141 za80g11.s 675 yo23d06.r 36341 BB63634 851 yo07f03.r 662 yo12e12.r	1909448 603065 424936 tg19b08 424936 tg19b08 807913 wf52c03 140043 qa8PaoV M783972 K-ESTO M783979 K-ESTO M740106 K-ESTO M740106 K-ESTO M76572 PM4-CT 700459 wd14b11 41522 yn90f12 70143 za80h09 M826648 K-ESTO 934992 op33h05 46211 y014h09 46211 y014h09 46211 g014h09	cription

## ALIGNMENTS

RESULT 1
BI909448
BI910000
BI90000
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KEYWORDS
SOURCE
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ORGANISM
ESTED
AUTHORS
NIH
MANU

BI909448 767 bp mRNA linear EST 16-OCT-2001 603065172F1 NIH\_MGC\_118 Homo sapiens cDNA clone IMAGE:5214568 5', mRNA sequence.
BI909448

BI909448.1 GI:16172666 EST. human.

Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 767)

1 (bases 1 to 767)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)

em\_gss\_rod:\*

EST 03-JAN-2000

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US-09-993-234-6_COPY_25_198 (1-174) x BI909448 (1-767)
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                   CTGCCTGCCTGCTTCTATGAACATGGCGATGGCTGCGTGTCCTGCCCCACGAGCACCCT
                                       rCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysProThrSerThrLe 160
                                                                                                               SGlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThrAspCysGlyTh 140
                                                                                                                                                                       UCYSG1nValSerG1nCysValSerSerSerProPheTyrCysG1nProCysLeuAspCy 120
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                                                                                                                                                                                                                                                                                                                                                   sHisAsnSerGluCysAlaArgCysGlnAlaCysAspGluGlnAlaSerGlnValAlaLe
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Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium
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Plate: LLAM11538 row: o column: 17
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Contact: Robert Strausberg, Ph.D.
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Clone distribution: MGC clone distribution information can be
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/clone="IMAGE:5214568"
/clone_lib="NIH_MGC_118"
/tissue_type="leukocyte"
/lab_host="DH108"
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                                                                             LeuPheCysCysArgGlyCysProAlaGlyHisTyrLeuLysAlaProCysThrGluPro
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                                                         CTGTTTTGTTGCAGAGGCTGCCCAGCGGGGCACTACCTGAAGGCCCCTTGCACGGAGCCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert Strausberg, Ph.D.
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632 bp mRNA linear EST 03-JAN-
xv41bl2.xl Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone
IMAGE:2815679 3', similar to SW:WSL1_HUMAN Q93038 WSL-1 PROTEIN
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Location/Qualifiers
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/clone_lib="Soares_NFL_T_GBC_S1"
/lab_host="DH10B"
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                   Tissue Procurement: Ash Alizadeh, John Byrd, M.D., Mike Greve M.D., Louis M. Staudt, M.D., Ph.D.
CDNA Library Preparation: M. Bento Soares, Ph.D.
CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing C Clone distribution: NCI-CGAP clone distribution information found through the I.M.A.G.E. Consortium/LLNL at:
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similar to
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Insert Length: 2028 Std Error: 0.00
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Email: cgapbs-r@mail.nih.gov
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National Cancer Institute, Cancer Genome Ana
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h quality sequence stop: 445.
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                                                  adaptors
                                                                                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="inAGE:2109207"
/clone=lib="NCI_CGAP_CLL1"
/clone_lib="NCI_CGAP_CLL1"
/tissue_type="B-cell, chronic lymphotic leukemia"
/lab_host="DH10B"
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TR:000276 000276 LYMPHOCYTE
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    Not I and Eco RI sites of rary is normalized, and was
double-stranded cDNA was ligated to Eco RI (Pharmacia), digested with Not I and cloned into I and Eco RI sites of the modified pT7T3 vector, is normalized, and was constructed by Bento
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Primates;
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CDNA clone IMAGE:2109207 3'
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Query Match:
                                                      COMMENT
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                                                                                                                                                                                                                                             AI807913 698 bp mRNA linear EST 19-DEC-
wf52c03.x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone
IMAGE:2359204 3' similar to SW:WSL1_HUMAN Q93038 WSL-1 PROTEIN
    Email: cgapbs-remail.nih.gov

Email: cgapbs-remail.nih.gov

This clone is available royalty-free through LLNL; contact the

This clone is available royalty-free through LLNL; contact the
                                                                                                                                                                                             EST
                                                                   Unpublished (1997)
                                                                                 National Cancer Institute, Tumor Gene Index
                                                                                                                        Mammalia; Eutheria;
1 (bases 1 to 698)
                                                                                                                                                                                                                         AI807913
                                                      Contact: Robert Strausberg, Ph.D.
                                                                                                          NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
                                                                                                                                                    Eukaryota; Metazoa;
                                                                                                                                                                 Homo sapiens
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Insert Length:
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(info@image.linl.gov)
                                                                                                                                     Chordata;
Primates;
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Matches:
Conservative:
Mismatches:
Indels:
                                                                                              Cancer
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DB:
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                                                     CysLeuProGlyPheTyrGluH1sGlyAspGlyCysValSerCysProThrSerThrLeu
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h quality sequence stop: 459
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note="Organ: pooled; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Equal amounts of plasmid DNA from three normalized libraries (fetal lung NbHL19W, testis NHT, and B-cell NCI_CGAP_GCB1) were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 297480-302087, 682632-687239, 726408-728711, and 729096-731399. Subtraction by Bento Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone_lib="Soares_NFL_T_GBC_S1"
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/db_xref="taxon:9606"
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97.70%
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Conservative:
Mismatches:
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US-09-993-234-6_COPY_25_198 (1-174) x AI140043 (1-562)
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Query Match:
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81 GluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGlyTrpPheValGlu
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Email: cgapbs-remail.nih.gov
This clone is available royalty-free through LLNL;
IMAGE Consortium (info@image.llnl.gov) for further
INAGE Consortium (info@image.llnl.gov) for further
Insert Length: 1640 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Insert Length: 1640 Std Error: 0.00 Seq primer: -40ml3 fwd. ET from Amersham High quality sequence stop: 494.
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National Cancer Institute, Cancer Genome Anat
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Mammalia; Eutheria;
1 (bases 1 to 562)
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/lab_host="DHIOB (ampicillin resistant)"
/note="Organ: heart; Vector: py773D (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; 1st strand cDNA was primed with a Not I - oligo(dT) primer [5' TGTTACCAATCTGAACTGGGACGGCCGAECTFTTTTTTTTTTTTTTTTTT 3'], double-stranded cDNA was size selected, ligated to Eco RI adapters (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified py773 vector (Pharmacia). Library went through one round of normalization to a Cot = 5. Library constructed by M.Fatima Bonaldo. This library was constructed from the feature as the feature in the seame feature as the seame feature as the seame feature in the seame feature in the seame feature as the seame feature in the seame feature as the seame feature in the seame feat
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/db_xref="taxon:9606"
/clone="IMAGE:1693710"
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Kim, N.S.,
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l: +82-42-860-4470
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/note-*Organ: Stomach; Vector: pCNS; Site_1: EcoRI; Site_2: NotI; The poly (A)+ RNA was dephosphorylated with bacterial alkaline phosphatase (BAP) and then decapped with tabacco acid pyrophosphatase (TAP). The decapped intact mRNA was ligated with DNA-RNA linker including EcoR I site by treatment of T4 RNA ligase and the first strand cDNA was synthesized from oligo dT-selected mRNA by priming with dT-tailed vector. The dT-tailed vector was adjusted to have about 60nt. The cDNA vector was circularized with E. coli DNA ligase after digestion of EcoRI which site is also included in vector. An RNA strand converted to a DNA strand by Okayama-Berg method. The obtained cDNA vectors were used for transformation of the cDNA libraries constructed by this method are full-length enriched cDNA library."
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/cell_type="Scattering
/cell_line="SNU-620"
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/db_xref="taxon:9606"
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Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL;
IMAGE Consortium (info@image.llnl.gov) for further i
Possible reversed clone: similarity on wrong strand
Possible reversed clone: polyT not found
                                                                                          Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
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National Cancer Institute, Cancer Genome Anatomy
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K-EST0017828 S6SNU620 Homo mRNA sequence.
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US-09-993-234-6_COPY_25_198 (1-174) x BM744307 (1-565)
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CysGlyAsnSerThrCysLeuValCysProGlnAspThrPheLeuAlaTrpGluAsnHis
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Plate: 17 row: C column: 06
High quality sequence stop: 565.
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Kim, N.S., Hahn, Y., (
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/cell_type="Soattering floating"
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Korea Research Institute of Bioscience & Biotechnology
52 Boeun-dong Yuseong-gu, Daejeon 305-333, South Korea
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Kim,N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim,M.R., Oh,K.J., Cheong,J.E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,S. and Kim,Y.S.
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Plate: 27 row: G column: 03
High quality sequence stop: 514.
Location/Qualifiers
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+82-42-860-4409
/note-*Organ: Stomach; Vector: pCNS; Site_1: ECORI; Site_2: NOtI; The poly (A)+ RNA was dephosphorylated with bacterial alkaline phosphatase (BAP) and then decapped with tabacco acid pyrophosphatase (TAP). The decapped intact mRNA was ligated with DNA-RNA linker including EcoR I site by treatment of T4 RNA ligase and the first strand cDNA was synthesized from oligo dT-selected mRNA by priming with dT-tailed vector. The dT-tailed vector was adjusted to have about 60nt. The cDNA vector was circularized with E. coli DNA ligase after digestion of EcoRI which site is also included in vector. An RNA strand
                                                                                                                                                                      /cell_type="Scattering floating"
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Korea Research Institute of Bioscience & Biotechnology 52 Eoeun-dong Yuseong-gu, Daejeon 305-333, South Korea Tel: +82-42-860-4470
                                                                                                            Eukaryota; Metazoa; Chordata;
Mammalla; Eutheria; Primates;
1 (bases 1 to 514)
Kim,N.S., Hahn,Y., Oh,J.H., Lo
Oh,K.J., Cheong,J.E., Sohn,H.)
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BM783979
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E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim
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Plate: 32 row: A column: 05
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/db_xref="taxon:9606"
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Email: yongsung@mail.kribb.re.kr
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Oh, K.J., Cheong, J.E., Sohn, H.Y., Kim, J.M., Park, H.S., Kim, S.
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                                                                                                                                                                                                                                                                           This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Insert Length: 1699 Std Error: 0.00 Seq.primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                                           Tumor Gene Index Unpublished (1997)
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National Cancer Institute, Cancer Genome Ana
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                                                                                                                                                                                                                                                                                                                                 Email: cgapbs-r@mail.nih
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          /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:1744827"
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/dev_stage="19 weeks"
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                                      Genome Research Center
Korea Research Institute of Bioscience
52 Boeun-dong Yuseong-gu, Daejeon 305-3
Tel: +82-42-860-4470
Fax: +82-42-860-4409
                                                                                                                    Kim, Y.S.
21C Frontier Korean
Unpublished (2002)
                                                                                                                                                               Kim, N.S., Hahn, Y., Oh, J.H.,
Oh, K.J., Cheong, J.E., Sohn,
                                                                                                                                                                            Mammalia; Eutheria;
1 (bases 1 to 488)
Kim, N.S., Hahn, Y.,
                                                                                                                                                                                                                                                                                                      BM770798
K-EST0054419 S6SNU620s1
5', mRNA sequence.
            Email: yongsung@mail.kribb.re.kr
Plate: 21 row: D column: 12
                                                                                                           Contact: Kim YS
                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                               Ahn,H.Y., Chu,M.Y.,
J.M., Park,H.S., Kim,
                                                                   & Biotechnology
333, South Korea
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 with tabacco acid pyrophosphatase (PAP). The decapped intact mRNA was ligated with DNA-RNA linker including Ecor I site by treatment of T4 RNA ligase and the first strand cDNA was synthesized from oligo dT-selected mRNA by priming with dT-tailed vector. The dT-tailed vector was circularized with E. coli DNA ligase after digestion of EcoRI which site is also included in vector. An RNA strand converted to a DNA strand by Okyama-Berg method. The obtained cDNA vectors were used for transformation of competent cells E. coli TOPALOT by electroporation method. The cDNA libraries constructed by this method are full-length enriched cDNA library. After analyzing and sequencing about 2,000 - 3,000 colonies in original cDNA library, the abundant cDNAs were selected and amplified by PCR reaction using vector region primer including T7 promotor as 5' primer and N(dT)14 as 3' primer. The PCR products were used as template for synthesis of biotinylated single stranded RNA probes were hybridized with antisense single stranded cDNAs propared from original liberary and incubated with avidin-gel. After removing libraries were constructed by transformation of the remaining DNA into competent cells E. coli ToplOF' with electroporation method."

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/clone_lib="S6SNU620s1"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information car
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM.1833 row: n column: 03
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC) Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AGENCOURT_6794093 NIH_MGC_121
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5770562"
/clone_lib="NIH_MGC_121"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                              directionally cloned (ECORV site is destroyed upon cloning). Average insert size 1.7 kb, insert size range 0.7-3.5 kb. Library is normalized and enriched for full-length clones and was constructed by C. Gruber (Invitrogen). Research Genetics tracking code 017. Note this is a NIH_MGC Library."
                                                                                                                                                                                                                                                                                                                                                                                                         /note="Organ: brain; Vector: pCMV-SPORT6; Site_1: NotI; Site_2: EcoRV (destroyed); RNA source anonymous pool of fetal brains, female age 20 weeks, female age 24 weeks, and male age 26 weeks. Library is oligo-dr primed and
                                                                                                                 8.46e-73
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100.00%
100.00%
83.04%
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1 Homo sapiens cDNA clone IMAGE:5770562
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                                                                                                                                                                                                                                                                                                                                                                                                                                           AW002222 478 bp mrNA linear EST 09-MAR-2000 wt85c09.x1 Soares_thymus_NHFTh Homo sapiens cDNA clone IMAGE:2514256 3' similar to TR:014866 014866 SOLUBLE DEATH RECEPTOR
                                                                                                                                                                                                                                         1 (bases 1 to 478)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
Tumor Gene Index
Unpublished (1997)
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Mammalia; Eutheria;
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Location/Qualifiers
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Primates;
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Search completed: April Job time: 1093.39 secs

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                                                      GlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThrAspCysGlyThr
                                                                                                                                                   GluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGlyTrpPheValGlu
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                                                                                                                                                                                                                                           CysGlyAsnSerThrCysLeuValCysProGlnAspThrPheLeuAlaTrpGluAsnHis
                                                                                                                                                                                                                                                                              CTGTTTTGTTGCAGAGGCTGCCCAGCGGGGCACTACCTGAAGGCCCCTTGCACGGAGCCC
                                            GGGGCCCTGCACCACAA-CGGCTACTCTGTTC-CGCAGAGATCT-GACTGTGGGACC
                                                                                         TGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGCCAACCATGCCTAGACTGC
                                                                                                                                       GAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCCAGGCTGGTTTGTGGAG
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Result
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MODEL-frame+_p2n.model -DEV-xlh

-Q-ggn2_1/USPTO_Spool_VG0993234/runat_27032003_115457_15378/app_query.fasta_1.2346

-Q-fgn2_1/USPTO_Spool_VG09993234/runat_27032003_115457_15378/app_query.fasta_1.2346

-DB-Issued_Patents_NA -QFMT-fastap -SUFFIX=rn1 -MINMATCH-0.1 -LOOPCL-0

-LOOPEXT-0 -UNITS-blts -START-1 -END=-1 -MATRIX=blosum62 -TRANS-human40.cd1

-LIST-45 -DOCALLIGN-200 -THR_SCORE-pct -THR_MAX=100 -THR_NIN-0 -ALIGN-15

-MODE-LOCAL -QUTFMT-pto -NORM-ext -HEAPSIZE=500 -MINLEN-0 -MAXLEN-200000000

-MODE-LOCAL -QUTFMT-pto -NORM-ext -HEAPSIZE=500 -MINLEN-0 -MAXLEN-2000000000

-USER-US0993234_8CGN _1_1116_8runat_27032003_115457_15378 -NCPU-6 -ICPU-3

-NO_MARP -LARGEQUERY -NEG_SCORES-0 -WAIT -LONGLOG -DEV_TIMEOUT=120

-WARN TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPEXT=7

-YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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Xgapop 10.0 , X
Ygapop 10.0 , Y
Fgapop 6.0 , F
Delop 6.0 , F
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length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                               /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/BCOMB.seq:*
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Fgapext
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US-08-928-609-11
US-08-828-683A-9
US-08-928-609-5
US-08-928-609-2
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US-08-928-603A-2
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## ALIGNMENTS

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RESULT 1
US-08-815-469-3
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                                                                        COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/815,469
FILING DATE: HEREWITH
CLASSIFICATION: 435
PRIOR APPLICATION BATA:
APPLICATION NUMBER: No. 6153402 Yet Assigned
FILING DATE: 06-FEB-1997
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Yu, Guo-Liang
APPLICANT: Ni, Jian
APPLICANT: Dixit, Vishva
APPLICANT: Gentz, Reiner L.
APPLICANT: Dillon, Patrick J.
TITLE OF INVENTION: Death Domai
                   APPLICATION NUMBER: US 60/028,711 FILING DATE: 17-OCT-1996 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
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ZIP: 200
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1100 New York Ave.,
NUMBER:
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US 60/013,285
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Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 1488
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
TELEPAX: 202-371-2540
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 1254 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: both
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FEATURE:
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LOCATION:
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REGISTRATION NUMBER:
                                      ArgHisCysTrpProHisLysProLeuValThrAlaAspGluAlaGlyMetGluAlaLeu
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                                                                                                                  GlyAlaLeuHisArgHisThrArgLeuLeuCysSorArgArgAspThrAspCysGlyThr
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Mismatches:
Indels:
Gaps:
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                                   APPLICATION NUMBER: 60/026943
FILING DATE: 09/23/1996
ATTORNEY/AGENT INFORMATION:
NAME: MAISCHANG, Diane L.
REGISTRATION NUMBER: 5,600
REFERENCE/DOCKET NUMBER: 91052R1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5116
            TELEPHONE: 650/225-5416
TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 1
                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Ashken
                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                         APPLICATION NUMBER: US/08/928,069
FILING DATE: 11-Sep-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Ashkenazi, Avi
TITLE OF INVENTION: Apo-3
NUMBER OF SEQUENCES: 15
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STREET: 1 DNA Way
CITY: South San Fra
STATE: California
COUNTRY: USA
ZIP: 94080
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  CHARACTERISTICS:
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DB:
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Best Local Similarity:
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STRANDEDNESS: Single
TOPOLOGY: Linear
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SerArgAlaLeuGlyProAlaAlaAlaProThrLeuSerProGluSerProAlaGlySer 300
                                 TyrProGluThrGlnGluAlaLeuCysProGlnValThrTrpSerTrpAspGlnLeuPro
                                                                           ProAspSerSerGluLysIleCysThrValGlnLeuValGlyAsnSerTrpThrProGly
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                     TACCCCGAGACCCAGGAGGCGCTCTGCCCGCAGGTGACATGGTCCTGGGACCAGTTGCCC
                                                                CCTGACAGCAGTGAGAAGATCTGCACCGTCCAGTTGGTGGGTAACAGCTGGACCCCTGGC
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Alignment Scores: Pred. No.: Score:
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GENERAL INFORMATION:
                                                                                                                                                              INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 1634 base pairs
TYPE: Nucleic Acid
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FILING DATE: 31-Mar-1997
CLASSIFICATION: <UNKnown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/625328
FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
ATTORNEY/AGENT INFORMATION:
NAME: MAISCHARP, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007P1
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MEDIUM TYPE: 3.5 inch, 1.44 Mb f
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
TELEFAX: 650/952-9881
                                                                                                                TOPOLOGY: Linear SEQUENCE DESCRIPTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-2 LI AND Apo-3
NUMBER OF SEQUENCES: 28
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CITY: South San F
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Percent Similarity: Best Local Similarity:

Query Match DB:

1300 380 1240 360

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US-09-993-234-6_COPY_25_417 (1-393) x US-08-828-683A-9 (1-1634)
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                                                                   ArgArgTrpLysGluPheValArgThrLeuGlyLeuArgGluAlaGluIleGluAlaVal
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  Percent Similarity:
Best Local Similarity:
                                                   Alignment Scores:
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US-08-815-469-1
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 60/028,711
FILING DATE: 17-OCT-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/013,285
FILING DATE: 12-MAR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Steffe, Eric K.
REGISTRATION NUMBER: 36,688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1,
                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: No. 61
FILING DATE: 06-FEB-1997
PRIOR APPLICATION DATA:
                                          .
No.:
                                                                                                                  MOLECULE TYPE: FEATURE:
                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 14
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS:
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APPLICANT: Dillon, Patrick J.
TITLE OF INVENTION: Death Domain Containing
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                                                                                                                                       LENGTH: 1783 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPDLOGY: both
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LOCATION:
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1100 New York Ave.,
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Gentz, Reiner
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ValLeuLeuAlaGlyLeuValValProLeuLeuGlyAlaThrLeuThrTyrThrTyr
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                                    ProAlaMetMetLeuGlnProGlyProGlnLeuTyrAspValMetAspAlaValProAla
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                            CCAGCCATGATGCTGCAGCCGGGCCCGCAGCTCTACGACGTGATGGACGCGGTCCCAGCG
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                                                                                                                                                                Query Match:
                                                                                                                                                                                                       Score:
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Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb flopi
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/928,069
FILING DATE: 11-Sep-1997
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                      TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 1438 base pair
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TITLE OF INVENTION: Apo-3
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/0:
APPLICATION NUMBER: 60/0:
FILING DATE: 09/23/1996
ATTORNEY/AGENT INFORMATION:
NAME: ATTORNEY/AGENT DATE DATE OF
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STREET: South San Fra
CITY: South San Fra
CTATE: California
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REGISTRATION NUMBER: 35,
REFERENCE/DOCKET NUMBER:
                                                   21
                                                                                                                                                                                                                                                                    STRANDEDNESS:
TOPOLOGY: Lir
                                                                                                                                                                                                                                                                                              TYPE:
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                        LeuPheCysCysArgGlyCysProAlaGlyHisTyrLeuLysAlaProCysThrGluPro
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0. 6462176
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Indels:
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Matches:
Conservative:
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Alignment Scores:
                                              SEQUENCE DESCRIPTION: US-08-828-683A-5
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US-08-828-683A-5
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Patent No. 6469144
GENERAL INFORMATION:
APPLICANT: ASSKenazi, Avi
                                                                                                                               INFORMATION FOR SEQ ID NO: 5: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  161 GlySerCysProGluArgCysAlaAlaValCysGlyTrpArgGln 175
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NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007P1
                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/828,683A
FILING DATE: 31-Mar-1997
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Apo-2 LI AND Apo-3 POLYPEPTIDES NUMBER OF SEQUENCES: 28
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                                                                                                                                                                                                                                                             APPLICATION NUMBER: 08/625328
FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
                                                                                                                                                              TELEFAX: 650/952-9881
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                                                        SEQ ID NO:
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                                                 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/
FILING DATE: 11-Sep-1997
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                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Ashkena
          CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/
FILING DATE: 09/23/1996
                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb 1
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Apo-3
                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                         161 GlySerCysProGluArgCysAlaAlaValCysGlyTrpArgGln 175
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952.00
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Marschang, Diane

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US-08-828-683A-2
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APPLICANT: Ashkenazi, Avi J.
APILICANT: ASHKENAZION: Apo-2 LI AND
NUMBER OF SEQUENCES: 28
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REFERENCE/DOCKET NUMBER: P10
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
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TYPE: Nucleic Acid
STRANDEDNESS: Single
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb i
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Genente
                                                                                                                                                                                                                                                                                                                                      GlyAlaLeuHisArgHisThrArgLeuLeuCys-SerArgArgAsp 135
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                                                                                                    STATE: California COUNTRY: USA
                                                                                                                               CITY: South San Francisco
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US-08-828-683A-2
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US-08-762-308-10
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Query Match:
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                                                                                       Patent No. 5925548
GENERAL INFORMATION:
                                                                                                                       Sequence
                             APPLICANT: Beutler, APPLICANT: Bazzoni, TITLE OF INVENTION: ITILE OF INVENTION:
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SEQUENCE CHARACTERISTICS:
NUMBER OF SEQUENCES:
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REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: 91007P1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                  GlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHisLysLysIleGlyLeu
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                                                                                                                                                                                                                                                                                                               GAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCAGGGCTGGTTTGTGGAGTG
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                                                                                                       10, Application US/08762308
5. 5925548
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TYPE: Nucleic Acid
STRANDEDNESS: Single
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FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
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                                                                           Beutler, Bruce A.
ADDRESS
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Best Local Similarity:
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FILING DATE: 09-DEC-1996
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/224,593
FILING DATE: 05-APR-1994
ATTORNEY/AGENT INFORMATION:
NAME: Kitchell, Barbara S.
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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REFERENCE/DOCKET NUMBER: UT
TELECOMMUNICATION INFORMATION:
TELEPHONE: 418-3000
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CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
                                                       540
140 ThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysProThrSerThr 159
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                                                TGC-----TTCAACGGCACCGTGACAATCCCCTGTAAGGAGACTCAGAACACCGTGTGT
                                                                                CysGlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThrAspCysGly
                                                                                                                  GAGAACCAGTTCCAACGCTACCTGAGTGAGACACACTTCCAGTGCGTGGACTGCAGCCCC
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                                                                                                                                                                                                                                                        TACCTCAGG----CAGTGTCTCAGTTGCAAGACATGTCGGAAAGAAATGTCCCAGGTGGAG
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                                                    GENERAL INFORMATION:
APPLICANT: Taylor, J. Michael
APPLICANT: Kehrli, Jr., Marcus
APPLICANT: Lee, Eun-Kyung
                                                                                                                                            Sequence 1, Application US/09513007 Patent No. 6406907
  APPLICANT: Lee, Eun-KyundapplicanT: Mwangi, Simon TITLE OF INVENTION: BOVINITITLE OF INVENTION: AND N
                                                                                                                                                                                                                                                 1419
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                                                                                                                                                                                                                                                                                   391 ArgGlyPro
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NECROSIS FACTOR RECEPTOR-1
OF USE
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
LENGTH: 2440
TYPE: DNA
ORGANISM: Bos taurus
FEATURE:
NAME/KEY: CDS
LOCATION: (294)...(1706)
US-09-513-007-1
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CURRENT APPLICATION NUMBER: US/09/513,007
CURRENT FILING DATE: 2000-02-25
PRIOR APPLICATION NUMBER: 60/122,156
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 8
1053 AAAGAGGGGGAGCCAGAACTCCTGGTCCCGGCCCCAGGCTTCAACCCCACCACCACCATC 1112
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                                                                                                                                                                                                                                                                                                                ValSerCysProThrSerThrLeuGlySerCysProGluArg---CysAlaAlaValCys 171
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                                                                                                         TyrArgHisCysTrpProHisLys---
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                                                                   TACCAGCGG----TGGAAGCCCCAAGCTCTACTCCATCATTTGCGGGCAGTCGACTCTGGTA 1052
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O8-050-319B-24  08-050-319B-24  equence 24, Application US/08050319B  atent NO. 5633145  APPLICANT: M.Feldmann, P.W. Gray, APPLICANT: M.Feldmann, P.W. Gray, APPLICANT: M.J.C. Turner, F.M Brennan TITLE OF INVENTION: Modified human TNFalpha (Tumor TITLE OF INVENTION: Mccrosis Factor alpha) Receptor NUMBER OF SEQUENCES: 57  CORRESPONDENCE ADDRESS: ADDRESSEE: Reed & Robbins STREET: 635 Bryant Street CITY: Palo Alto STATE: California COUNTAX: USA ZIP: 94301  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIN Release #1.0, version #1.25 CURRENT APPLICATION DATA: APPLICATION: 435 CLASSIFICATION: 435 ATTORNEY/AGENT INFORMATION: NAME: Robbins, Roberta L.	16		AlavalTyrAlaAlaLeuGluArgMetGlyLeuAspGlyCysValGluAspLeuArgSer::::	GlnTyrGluMetLeuLysArgTrpArgGlnGlnGlnProAlaGlyLeuGly:	LeuArgGluAlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGln:	TyrAspValMetAspAlaValProAlaArgArgTrpLysGluPheValArgThrLeuGly	LeuSerProGluSerProAlaGlySerProAlaMetMetLeuGlnProGlyProGlnLeu	ValThrTrpSerTrpAspGlnLeuProSerArgAlaLeuGlyProAlaAlaAlaAlaProThr :	GlyAsnSerTrpThrProGlyTyrProGluThrGlnGluAlaLeuCysProGln: ::::::	AlaHisThrLeuLeuAlaProProAspSerSerGluLysIleCysThrValGlnLeuVal :	GlualaGlyMetGlualaLeuThrProProProAlaThrHisLeuSerProLeuAspSer:		lThrAlaAsp
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Best Local Similarity:
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REFERENCE/DOCKET NUMBER: 5150-0030
RELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 617-8999
TELEPHAX: (415) 327-3231
INFORMATION FOR SEQ ID NO: 24:
                                                 608
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LENGTH: 2062 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
203 CysTrpProHisLys---
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               257 GGGGACAGGGAGAGAGAGATAGTGTGTGTCCCCAAGGAAAATATATCCCACCCTCAAAAT 316
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                                       TICTITGGTCTTTGCCTTTTATCCCTCCTCTTCATTGGTTTAATGTATCGCTACCAACGG 868
                                                                     LeuAlaGlyLeuValValProLeuLeuLeuGlyAlaThrLeuThrTyrThrTyrArgHis
                                                                                                                                                                                                    ThrSerThrLeuGlySerCysProGlu-----ArgCysAlaAlaValCys-----
                                                                                                                                                                                                                                        ACCGTGTGCACCTGCCATGCAGGTTTCTTTCTAAGAGAAAACGAGTGTGTCTCCTGT---
                                                                                                       ATTGAGAATGTTAAGGGCACTGAGGACTCAGGCACCACAGTGCTGTTGCCCCCTGGTCATT
                                                                                                                                                                       ------AGTAACTGTAAGAAAAGCCTGGAGTGCACGAAGTTGTGCCTACCCCAG
                                                                                                                                                                                                                                                                     AspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysPro 156
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USESCO CONTRACTOR CONT	dd Vy	₽ 5	P B	Qy	DP OA	dd	Ð	δ <u>β</u>	γ <sub>2</sub>	B 5	DB DB	Qy	Db Qy	문	Qy	Dβ
S S	Qy 387 SerArgLeu 389       Db 1469 GAGGCGCTT 1477	14	1349	350	Qy 331 GlyLeuArgGluAlaGluIleGluAlaValGluIleGlyArgPheArgAsp        :::      :::      :::           1289 GGGCTGAGCGACCACGAGATCGATCGGCTGGAGAGCGGGGGGGG	, <del></del>	Db 1169	рр 1109 Qy 295		Db 1064 TGTCCCAACTTTGCGGCTCCCGGCAGAGAGGGTGGCACCACCC	1037	249	OY 234AlaHisThrLeuLeuAlaProProAspSerSerGluLysIleCys  Db 977 CCCACTCCAGGCTTCACCCCCACCCTGGGCTTCAGTCCCAGTTCCACCTTCACC	926 GAGCTTGAAGGAACT	216 GlyMetGluAlaLeuThrProProProAlaThrH :::	Db 869TGGAAGTCCAAGCTCTACTCCATTGTTTGTGGGAAATCGACACCTGAAAAAAGAGGGG
		<u>μ</u> ω		ω .	<u></u>	μ ω	<b>–</b> (	TI	SerPro 2	ro    CACCCTAT				AGCTTCAGT 9	Ser 2	AAAGAGGGG 9
		86 468	4	99	34.9	288	N F		94	286 1108	1063	268	248	976	233	925

ATTORNEY/AGENT INFORMATION: NAME: Robbins, Roberta L

Robbins, Roberta L.

APPLICATION NUMBER: US/O FILING DATE: 10-May-1993

US/08/050,319

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DB:
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TELEFAX: (415) 327-323:
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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REFERENCE/DOCKET NUMBER: 51:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 617-8999
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FEATURE:
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LOCATION:
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STRANDEDNESS: double
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LeuAlaGlyLeuValValProLeuLeuGlyAlaThrLeuThrTyrThrTyrArgHis 202
                                                                                                                                                                                                                                                                     CysLeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThr 136
                                                                                                                                                                                                                                                                                                      CATTATTGGAGTGAAAACCTTTTCCAGTGC-----TTCAATTGCAGCCTC 598
                                              ATTGAGAATGTTAAGGGCACTGAGGACTCAGGCACCACAGTGCTGTTGCCCCTGGTCATT
                                                                                                                                                                                                       AspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysPro
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                                                                                                              -----AGTAACTGTAAGAAAAGCCTGGAGTGCACGAAGTTGTGCCTACCCCAG
                                                                                                                                           ThrSerThrLeuGlySerCysProGlu-----ArgCysAlaAlaValCys-----
                                                                                                                                                                         ACCGTGTGCACCTGCCATGCAGGTTTCTTTCTAAGAGAAAACGAGTGTGTCTCCTGT---
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                                                                                                                                                                                                                                     Sequence 1, Application US/09106038A
Patent NO. 6007995
GENERAL INFORMATION:
APPLICANT: Brenda F. Baker and Le
TITLE OF INVENTION: ANTISENSE MOI
TITLE OF INVENTION: EXPRESSION
             MEDIUM TYPE: 3.5 inch disk, 1.44
COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows NT
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inc
                                                                                                                                                                                                                                                                                                                                                                                       1469 GAGGCGCTT
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                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
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                                                                                                              ZIP:
 APPLICATION NUMBER:
                                                                                                                             COUNTRY:
                                                                                                                                                STATE:
                                                                                                                                                             CITY: Carlsbad
                                                                                                                                                                           ADDRESSEE: Isis Pharmaceuticals, STREET: 2292 Faraday Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                     SerArgLeu 389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GlyLeuArgGluAlaGluIleGluAlaValGluValGluIleGlyArg---PheArgAsp 349
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CAGAAGTGGGAGGACAGTGCCCACAAGCCACAGAGCCTAGACACTGATGACCCCCGGACG
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                                                                                                                                                                                                                             EXPRESSION
91
US/09/106,038A
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Best Local Similarity:
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CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Laurel Spear Bernstein
REGISTRATION NUMBER: 37,280
REFERENCE/DOCKET NUMBER: RTS-0004
TELECOMMUNICATION INFORMATION:
TELEPHONE: (760) 931-9200
TELEPAX: (760) 603-3820
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 2161
TPUED: NUMBER: 3821
                                   910 TTCTTTGGTCTTTGCCTTTATCCCTCCTCTTCATTGGTTTAATGTATCGCTACCAACGG 969
                                                                      183 LeuAlaGlyLeuValValProLeuLeuLeuGlyAlaThrLeuThrTyrThrTyrArgHis 202
                                                                                                                                                                                                                         157 ThrSerThrLeuGlySerCysProGlu-----ArgCysAlaAlaValCys------ 171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      595
         203 CysTrpProHisLys--
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               538 CACCTCAGA---CACTGCCTCAGCTGCTCCAAATGCCGAAAGGAAATGGGTCAGGTGGAG 594
                                                                                                                                                                                                                                                                                                                                                                                                                                         100 GluCys-----GlnValSerGlnCysValSerSerSerProPheTyrCysGlnPro 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       478 CCGGGGCAGGATACGGACTGCAGGGGGTGTGAGAGCGGCTCCTTCACCGCTTCAGAAAAC 537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              418 AATTCGATTTGCTGTACCAAGTGCCACAAAGGAACCTACTTGTACAATGACTGTCCAGGC 477
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                60 HisHisAsnSerGluCysAlaArgCysGlnAlaCysAspGluGlnAlaSerGlnValAla 79
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STRANDEDNESS: sing
TOPOLOGY: linear
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                                                                                                           ATTGAGAATGTTAAGGGCACTGAGGACTCAGGCACCACAGTGCTGTTGCCCCCTGGTCATT 909
                                                                                                                                               ------GlyTrpArgGlnMetPheTrpValGlnValLeu 182
                                                                                                                                                                                        ------AGTAACTGTAAGAAAAGCCTGGAGTGCACGAAGTTGTGCCCTACCCCAG
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                                                                                                                                                                                                                                                                                                  AspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysPro 156
                                                                                                                                                                                                                                                                                                                                       TGCCTCAAT---GGGACCGTGCAC------CTCTCCTGCCAGGAGAAACAGAAC 744
                                                                                                                                                                                                                                                                                                                                                                          CysLeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThr 136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATCTCTTCTTGCACAGTGGACCGGGACACCGTGTGTGGCTGCAGGAAGAACCAGTACCGG 654
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369.00
39.05%
28.44%
16.70%
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Matches:
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Alignment Pred. No.: Score: Percent Si	US-01	α	, , , , , , , , , , , , , , , , , , ,	RESULT US-09- ; Sequ ; Pate	ф	Qy	Db	Qy b	Qy Dh	Db	Оу	pb :	Qy g	P 27	Db	Оу	Db	Ov	Db Qy	Db	Оу	Db	VO V	dd://
ment Scores: No.: nt Similarity:	0 Z K Z Z	AR O	L INFO	ESULT 14 3-09-505-250-3 Sequence 3, Applicati Patent No. 6329148	1570 GAGGCGCTT	387 SerArgLeu	0	367 GlyAlaVal	350	1390 GGGCTGAGC	331 GlyLeuArç		311 LeuTyrAsı	295	1210 CAGGGGGC	287	5 -	φ	249 ThrValG1 ::: 1138 TCC		234	1027 GAGCTIGA	216 GlyMetGl	970TGGAA
2.2e-21 369.00 39.05%	piens (1623)	NUMBER: US/0 2000-02-15 : 4 or Windows V	RMATION: ROSen, Glenn ROSen, Glenn RAO, Peter VENTION: Synergistic VENTION: Triptclides NCE: SUN-109PRV2	on US/095052	1578	389	GACGCGTGCTCCGCG	AGCATGCTGGCGACCT TyrAlaAlaLeuGlua	lnGlnTyrGluMetLeuLysArgTrpArgGlnGlnGlnPro	GACCACGAGATCGATO	GlyLeuArgGluAlaGluIleGluAlaValGluValGluIleGlyArg	: : : : : :	3GAGGACAGCGCCCACI ValmetAspAlaVali	GluSerProAlaGly	GACCCCATCCTTGCG	! ! ! ! ! ! ! !		ValThrTrnserTrn	nLeuValGlyAsnSer :::::::	AGGCTTCACCCCCACC	AlaHisThr	GAGCTTGAAGGAACTACTACTAAGCCC-	lyMetGluAlaLeuThrProProProAlaThrHisLeuSerProLeuAspSer	GTCCAAGCTCTACTCC
Length: Matches: Conservative		9/505,250A ersion 3.0	Anti-Cancer Th	50A			Hecety Ledaspe	GGAGGCGCGCACGC	TpArgGlnGlnGlnP	:::   :::::: GGCTGGAGCTGCAGA	AlaValGluValGluI	CCCCGTTGCGCTGGA	AAGCCACAGAGCCTAG	GluSerProAlaGlySerProAlaMetMetLeuGlnProGlyPro-		AlaAlaA		AspClpTouBroCor.	TrpThrProGlyTyrE :::	CTGGGCTTCAGTCCC	LeuLeuAlaProPro		ProAlaThrHisLeu	ATTGTTGTGGGAAA
2161 126 : 47			erapy with in Ligands				AGCTGCTGGGACGCGTGCTCCGCGACATGGACCTGGTGGGCTGCCTGGAGGACATCGAG	CGCGGCGCGAGGCCACGCT	GlnGlnTyrGluMetLeuLysArgTrpArgGlnGlnGlnProAlaGlyLeu		leGlyArgPheArgAs	CTGTACGCCGTGGTGGAGAACGTGCCCCCGTTGCGCTGGAAGGAA	AWAAWIYGGAGGACAGCCCCACAAGCCACAGAGCCTAGACACTGATGACCCCGCGACG euTyrAspValMetAspAlaValbroalaArgargTrotucclubboValaargacct	euGlnProGlyProGln 		AlaAlaAlaProThrLeuSerPro	LeurioseiArgAlaLeuGlyPro		ThrValGlnLeuValGlyAsnSerTrpThrProGlyTyrProGluThrGlnGluAlaLeu :::: TCCAGGTCACTTATATATATATATATATATATATATATATATA	CCCACTCCAGGCTTCACCCCCACCCTGGGCTTCAGTCCCGTGCCCAGTTCCACCTTCACC	AlaHisThrLeuLeuAlaProProAspSerSerGluLysIleCys	TTCA	SerProLeuAspSer	TGGAAGTCCAAGCTCTACTCCATTGTTTGTGGGAAATCGACACCTGAAAAAGAGGGG
							9 386 G 1569	15	u 366			и 330   A 1389		n 310			7 1209		268	C 1137			233	GG 1026

294 1269	AlaAlaAlaProThrLeuSerPro	287 1210	Db Qy
286 1209	CysProGlnValThrTrpSerTrpAspGlnLeuProSerArgAlaLeuGlyPro	269 1165	Db Qq
268 1164	ThrValGlnLeuValGlyAsnSerTrpThrProGlyTyrProGluThrGlnGluAlaLeu ::: TCCAGCTCCACCTATACCCCCGGTGAC	249 1138	Db Qq
248 1137	AlaHisThrLeuLeuAlaProProAspSerSerGluLysIleCys	234 1078	B 6
233 1077	GlyMetGluAlaLeuThrProProProAlaThrHisLeuSerProLeuAspSer :::	216 1027	В <b>о</b>
215 1026	CysTrpProHisLysProLeuValThralaAspGluAla	203 970	Db Qy
202 969	LeuAlaGlyLeuValValProLeuLeuGlyAlaThrLeuThrTyrThrTyrArgHis	183 910	D Oy
182 909		172 850	ДV
171 849	urLeuGly	157 802	Db Qq
156 801	AspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysPro	137 745	P 64
136 744	CysLeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThr       ::     :::              GCCTCAATGGGACCGTGCAC	117 700	g 6
116 699	GluCysGlnValSerGlnCysValSerSerSerProPheTyrCysGlnPro :::	100 655	D Q
99 654	LeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGlyTrpPheVal::::::   ::::::::::::::::::::::::::::	595	pb dg
79 594	HisHisAsnSerGluCysAlaArgCysGlnAlaCysAspGluGlnAlaSerGlnValAla	60 538	Db Qq
59 537	ProCysGlyAsnSerThrCysLeuValCysProGlnAspThrPheLeuAlaTrpGluAsn	40 478	g dy
39 477	GlyLeuPheCysCysArgClyCysProAlaGlyHisTyrLeuLysAlaProCysThrGlu	20 418	B 8
19 417	GlyThrArgSerProArgCysAspCysAlaGlyAspPheHisLysLysIle	358	DP 6
	7 (1-393) x US-09-505-250-3	993	US-
	Local Similarity: 28.44% Mismatches: 178 92 Match: 16.70% Indels: 92 Gaps: 21	t Loca ry Mat	Best I Query DB:

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US-08-321-668-1
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                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/321,668
FILING DATE: 12-OCT-1994
CLASCIFICATION 12-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: WALLACH,
APPLICANT: BRAKEBUSC
                                                                            INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 2175 base pairs
TYPE: nucleic acid
                                                                                                                                                                                      NAME: BROWDY, ROGET L.
REGISTAN NUMBER: 25,618
REFERENCE/DOCKET NUMBER: WAN
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
                                                                                                                                                                                                                                                            CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: IL 107268
FILING DATE: 12-OCT-1993
ATTORNEY/AGENT INFORMATION:
NAME: BROWDY, ROGET L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: BRAKEDUSCH, CORD
APPLICANT: VARFOLOMEEV, Eugene
APPLICANT: BATKIN, Michael
TITLE OF INVENTION: THE THE RECEPTORS, THEIR PREPARATION AND THEIR USE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: BROWDY AND NEIMARK
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                               TOPOLOGY: 11
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                                                                                                                                            TELEFAX: 248633
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                                                               STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Washington
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NAME/KEY: CDS
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Percent Similarity:
Best Local Similarity:
Query Match:
DB:
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US-08-321-668-1
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                                                                                                                       CysLeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThr 136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CATTATTGGAGTGAAAACCTTTTCCAGTGC------TTCAATTGCAGCCTC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GlyLeuPheCysCysArgGlyCysProAlaGlyHisTyrLeuLysAlaProCysThrGlu
                             ThrValGlnLeuValGlyAsnSerTrpThrProGlyTyrProGluThrGlnGluAlaLeu 268
                                                              CCCACTCCAGGCTTCACCCCCACCCTGGGCTTCAGTCCCGTGCCCAGTTCCACCTTCACC 113
                                                                                                                                                                                    ---TGGAAGTCCAAGCTCTACTCCATTGTTGTGGGAAATCGACACCTGAAAAAAGAGGGG
                                                                                                                                                                                                                   CysTrpProHisLys-----
                                                                                                                                                                                                                                                                                                              ATTGAGAATGTTAAGGGCACTGAGGACTCAGGCACCACAGTGCTGTTGCCCCCTGGTCATT 909
                                                                                                                                                                                                                                                                                                                                                                                                                                        ACCGTGTGCACCTGCCATGCAGGTTTCTTTCTAAGAGAAAACGAGTGTGTCTCCTGT---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGlyTrpPheVal ::: :::|||::: :::
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 TCC-----AGCTCCACCTATACCCCCGGTGAC--
                                                                                 -----AlaHisThrLeuLeuAlaProProAspSerSerGluLysIleCys 248
                                                                                                                                                                                                                                                                                                                                                                         -----AGTAACTGTAAGAAAAGCCTGGAGTGCACGAAGTTGTGCCTACCCCAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   269 CysProGlnValThrTrpSerTrpAspGlnLeuProSerArgAlaLeuGlyPro-----
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GAGGCGCTT 1578
                                                                              GlyAlaValTyrAlaAlaLeuGluArgMetGlyLeuAspGlyCysValGluAspLeuArg
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                            SerArgLeu 389
                                                            GAGCTGCTGGGACGCGTGCTCCGCGACATGGACCTGCTGGGGCTGCCTGGAGGACATCGAG 1569
                                                                                                                         -----GluSerProAlaGlySerProAlaMetMetLeuGlnProGlyPro---Gln
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Search completed: April Job time: 87.3429 secs 6, 2003, 23:29:40

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-DB-(enEmbl -QFMT-fastap -SUFFIX=rge -MINMATCH=0.1 -LOOPCL=0 -LOOPEXT=0
-UNITS-bits -START=1 -END-1 -MATRIX=blosum62 -TRANS-human40.cd1 -LIST-45
-DOCALIGN=200 -THR_SCORE-pct -THR_MAX=100 -THR_MIN-0 -ALIGN=15 -MODE-LOCAL
-OUTPMT-pto -NORM-ext -HEARSIZE-500 -MINLEN-0 -MAXLEN-20000000
-USER-US09993234_eCGN_1_1_8534_erunat_27032003_115455_15349 -NCPU-6 -ICPU-3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPOEXT=0.5 -FGAPOP=6 -FGAPEXT=7
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-MODEL=frame+_p2n.model -DEV=xlh
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Yu,G.L., Ni,J., Gentz,R.L. anu
Peath domain containing receptors
Patent: US 6153402-A 3 28-NOV-2000;
Location/Qualifiers
1. .1254
2. cm="unknown"
226
                              CGCCACTGCTGGCCTCACAAGCCCCTGGTTACTGCAGATGAAGCTGGGATGGAGGCTCTG
        ArgHisCysTrpProHisLysProLeuValThrAlaAspGluAlaGlyMetGluAlaLeu
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RESULT 2
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Chinnalyan,A.M., O'Kourke,K., Yu,G.L., Lyons,R.H., G.Duan,D.R., Xing,L., Gentz,R., Ni,J. and Dixit,V.M. Signal transduction by DR3, a death domain-containing related to TNFR-1 and CD95
Science 274 (5289), 990-992 (1996)
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Chinnalyan,A.M., O'Rourke,K., Yu,G.-L., Lyons,R.
Duan,R., Xing,L., Gentz,R., Ni,J. and Dixit,V.M.
Direct Submission
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                                                                                /organism="Homo sapiens"
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	5 6 6	ProAspSerSerGluLysIleCysThrValGlnLeuValGlyAsnSerTrpThrProGl	79	B 6
	240 792	ThrProProProAlaThrHisLeuSerProLeuAspSerAlaHisThrLeuLeuAlaPro	y 221 b 733	å å
	220 732	ArgHisCysTrpProHisLysProLeuValThrAlaAspGluAlaGlyMetGluAlaLeu	y 201 b 673	g Qy
	200 672	ValLeuLeuAlaGlyLeuValValProLeuLeuGlyAlaThrLeuThrTyrThrTyr 	y 181 b 613	d Vo
	180 612	GlySerCysProGluArgCysAlaAlaValCysGlyTrpArgGlnMetPheTrpValGln	y 161 b 553	ρ δ
	160 552	CysLeuProGlyPheTyrGluH1sGlyAspGlyCysValSerCysProThrSerThrLeu 	y 141 b 493	g Qy
	140 492	GlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThrAspCysGlyThr 	y 121 b 433	B 6
	120 432	CysGlnValSerGlnCysValSerSerSerProPheTyrCysGlnProCysLeuAspCys 	y 101 o 373	B Oy
	100 372	GluasnCysSerAlaValalaaspThrargCysGlyCysLysProGlyTrpPheValGlu 	y 81 0 313	g VQ
	80 312	HisasnSerGluCysalaargCysGlnAlaCysAspGluGlnAlaSerGlnValAlaLeu 	y 61 5 253	g Q
	60 252	CysGlyAsnSerThrCysLeuValCysProGlnAspThrPheLeuAlaTrpGluAsnHis	y 41 b 193	B 6
	40 192	LeupheCysCysArgGlyCysProAlaGlyHisTyrLeuLysAlaProCysThrGluPro	y 21 b 133	8 6
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FEATURES Location/Qualifiers source 11254
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ACCESSION VERSION KEYWORDS SOURCE ORGANISM
RESULT 3 HSU78029 LOCUS DEFINITION
Qy 381 CysValGluAspLeuArgSerArgLeuGlnArgGlyPro
Qy 361 GlnGlnProAlaGlyLeuGlyAlaValTyrAlaAlaLeuGluArgMetGlyLeuAspGly
Qy 341 GluvaiGluIleGlyArgPheArgAspGlnGlnTyrGluMetLeuLysArgTrpArgGln
Qy 321 ArgArgTrpLysGluPheValArgThrLeuGlyLeuArgGluAlaGluIleGluAlaVal
Qy 301 ProAlaMetMetLeuGlnProGlyProGlnLeuTyrAspValMetAspAlaValProAla
Qy 281 SerargalaLeuGlyProAlaAlaAlaAroThrLeuSerProGluSerProAlaGlySer
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Ho
1 (bases 1 to 1254)
Screaton,G.R., Xu,X.N., Olsen,A.L., Cowper,A.E., Tan,R
McMichael,A.J. and Bell,J.I.
LARD: a new lymphoid-specific death domain containing
regulated by alternative pre-mRNA splicing
Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
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                                                                                                       /product="lymphocyte associated receptor of death la"
/protein_id="AACS1306.1"
/db_xref="g1:2071949"
/db_xref="g1:2071949"
/translation="MEORPRGCAAVAAALLLVLLGARAQGGTRSPRCDCAGDFHKKIG
LFCCRCCPAGHYLKAPCTEPGCNSTCLVCPQDTFLAWENHHNSECARCQACDEQASOV
ALENCSAVADTRCGCKPGWFVECQVSQCVSSPTVCQPCLUCGALHRHTRLLCSRRDT
DCGTCLDGFYERIGDGCVSCPTSTLGSCPERCAAVCGWRQMFWVQVLLAGLVVPLLLGA
TLTYTYRHCWPHKPLVTADEAGMEALTPPPATHLSPLDSAHTLLAPPDSSEKICTVOL
VGNSWTPGYPETQEALCPQVTWSWDQLDSRALGPAAPTLSPESPAGSPAMMLQPGPQ
LYDVNDAVPARWKETVTALGREAEIEAVEVEIGRFRDQQYEMLKRWRQQQPAGLGA
VYAALERMGLDGCVEDLRSRLGRGP"
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TNF-R1; contains a death
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/db_xref="taxon:9606"
/cell_line="HeLa"
                                                                                                                                                                                                                                                                                                                                              function="mediates apoptosis"
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         ArgArgTrpLysGluPheValArgThrLeuGlyLeuArgGluAlaGluIleGluAlaVal
                                                                          AGCAGAGCTCTTGGCCCCGCTGCTGCGCCCACACTCTCGCCAGAGTCCCCAGCCGGCTCG
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                                                                                                                                                                                                                                                                                                                                                                                                                              Goddard,A.D., Bauer,K.D. and Ashkenazi,A.
Apo-3, a new member of the tumor necrosis factor receptor contains a death domain and activates apoptosis and NF-kB Curr. Biol. (1996) In press
2 (bases 1 to 1634)
Marsters,S.A., Sheridan,J.P., Donahue,C.J., Pitti,R.M., Gr Goddard,A.D., Bauer,K.D. and Ashkenazi,A.
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Marsters, S.A., Sheridan, J.P., Donahue, C.J.,
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Apo-3 mRNA, complete cds.
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                                                                                                                               /note="contains death domain, NF-kB; TNF receptor family men
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/dev_stage="fetal"
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Ashkenazi,A.J., Baker
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                                     Mammalia; Eutheria; Chordata; Craniata; Vertebrata; Eutele Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1662)

RS Ashkenazi,A.J., Goddard,A., Godowski,P.J., Gurney,A.L., Hillan,K.J., Marsters,S.A., Pan,J., Pitti,R.M., Roy,M.A., Stone,D.M., Watanabe,C.K. and Wood,W.I.

Compositions and methods for the treatment of tumour Patent: WO 0153486-A 23 26-JUL-2001;

Genentech, Inc. (US)

Location:
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/organism="Homo sapiens"
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U83597
                   Direct Submission
Submitted (03-JAN-1997) Molecular Biotechnology, Univewashington, 1705 NE Pacific Street, HSB-K360, Seattle,
                                         Eukaryota; Metazoa; Chorda
Mammalia; Eutheria; Primat
1 (bases 1 to 1528)
Chaudhary, P.M. and Hood, L.
                                                                         Homo sapiens
                                                                                  Homo sapiens.
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Location/Qualifiers
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                                                          Chordata;
Primates;
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3 (DDR3) mRNA,
                                                        Craniata; Vertebrata; Catarrhini; Hominidae;
                                                                                                                                                   1481
                                                          Hominidae;
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                   University of attle, WA 98195,
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                            GlySerCysProGluArgCysAlaAlaValCysGlyTrpArgGlnMetPheTrpValGln 180
                                                                                     CysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysProThrSerThrLeu
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                GGGAGCTGTCCAGAGCGCTGTGCCGCTGTCTGTGGCTGGAGGCAGATGTTCTGGGTCCAG
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AVADTRCGCKPGWFVECQVSQCVSSSPFYCQPCLDCGALHRHTRLLCSRRDTDCGTCL
LGFYEHGDGCVSCPTSTLGSCPERCAAVCGWRQMFWQVLLAGLVVPLLLGATLTFYTY
RHCWPHKPLVTADEAGMEALTPPPATHLSPLDSAHTLLAPDSSEKICTVQLVGNSWT
PGYPETGBLCPQVTWSMDQLPSRALGFAAAPTLSPESPAGSPAMMLQPGPQLYDVM
AVPARWKEFVERIGELEAVENETENVEVEIGRFRDQQYEMLKRWRQQQPAGLGAVYAALE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /gene="DDR3"
/function="mediates apoptosis when cross-linked"
/note="A TWIR1-related death-domain containing receptor;
DR3, WSL-LR, Apo-3, TRAMP; transmembrane form; similar to
human EST clone 298913, GenBank Accession Number N71143"
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/db_xref="taxon:9606"
/chromosome="1"
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1 (bases 1 to 1557)

Bodmer,J.L., Burns,K., Schneider,P., Hofmann,K., Steiner,V., Thome,M., Bornand,T., Hahne,M., Schroter,M., Becker,K., Wilson,A., French,L.E., Browning,J.L., MacDonald,R. and Tschopp,J.

TRAMP, a novel apoptosis-mediating receptor with sequence homology to tumor necrosis factor receptor 1 and Fas(Apo-1/CD95)

Immunity 6 (1), 79-88 (1997)

9052839
                      Bodmer, J.L., Burns, K., Schneider, P., Hofmann, K., Steiner, V., Thome, M., Bornand, T., Hahne, M., Schroter, M., Wilson, A., French, L.E., Browning, J.L., MacDonald, R. and Tschopp, J. Direct Submission
Submitted (18-OCT-1996) Institute of Biochemistry, Universit Lausanne, 155 Chemin des Boveresses, Epalinges, CH 1066,
                Switzerland
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/product="apoptosis-mediating receptor TRAMP"
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NCSAVADITRGGCKPGNFVEQVSQCVSSSSFFVCQPCLDCGALHRHTRLLCSRRDTDCG
TCLLGFYEHGDGCVSCPTSTLGSCPERCAAVCGWRQMFWQVLLAGATUT
YTYRHCWPHKPLVTADEBAGMEALTPPPATHLSPLDSAHTLLAPDSSEKICTVQLVGN
SWTPGYPETQEALCPQVTWSWDQLPSRALGPAAPTLSPESPAGSPAMMLQPGPQLYD
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/map="1p36.2"
/tissue_type="fe
<1. .1245
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ALERWGLDGCVEDLRSRLQRGP"
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/organism="Homo sapiens"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                               Human lymphocyte associated receptor of death 1b m alternatively spliced, complete cds.

U94502
U94502.1 GI:2071950
                                               Screatin, G.R.

Direct Submission
Submitted (19-MAR-1997) Molecular Immunology
Molecular Medicine, John Radcliffe Hospital,
9DU, UK
                                                                                                                        Screaton,G.R., Xu,X.N., Olsen,A.L., Cowper,A.E., McMichael,A.J. and Bell,J.I.
LARD: a new lymphoid-specific death domain contaregulated by alternative pre-mRNA splicing Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619
                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 1257)
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/organism="Homo sapiens"
/db_xref="taxon:9606"
/cell_line="HeLa"
                             Location/Qualifiers
1. .1257
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                                    GGGAGCTGTCCAGAGCGCTGTGCCGCTGTCTGTGGCTGGAGGCAGATGTTCTGGGTCCAG
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Accession Number U94501, probably represents alternative
3' splice site"
421 c 408 g 226 t
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DCGTCLLPGFYEEGDGCVSCPTSTLGSCCPERCAAVCGWRQMFWYQVLLAGLVVPLLLGA
TLTYTTRHCWPHKPLVTAADEAGMEALTPPPATLSPLDSAHTLLAPPDSSEKICTVQ
LVGNSWTPGYVEETQEALCPQVTWSWDQLPSRALGPAAAPTLSPESPAGSPAMMLQPGP
QLYDVMDAVPARRWKEFVRTLGLREADELEAVEVEIGRFRDQQYEMLKRWRQQQPAGLG
AVYAALERWGLDGCVEDLRSRLQRGP*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /function="mediates apoptosis"
/note="LARD-1b; membrane protein;
TNF-R1; contains a death domain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /translation="meQRPRGCAAVAAALLLVLLGARAQGGTRSPRCDCAGDFHKKIG
LFCCRGCPAGHYLKAPCTEPCGNSTCLVCPQDTFLAWENHHNSECARCQACDEQASQV
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/protein_id="AAC51307.1"
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/organism="Homo sapiens"
/db_xref="taxon:9606"
/cell_line="HeLa"
/cell_type="lymphocyte"
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                                                                                                                                                                    Screaton,G.R., Xu,X.N., Olsen,A.L., Cowper,A.E., McMichael,A.J. and Bell,J.I.
LARD: a new lymphoid-specific death domain contaregulated by alternative pre-mRNA splicing proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619
                                Submitted (19-MAR-1997) Molecular I Molecular Medicine, John Radcliffe
                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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/protetn_id="AACS1308.1"
/protetn_id="AACS1308.1"
/db_xref="GI:2071953"
/tansilation="MEDRAPRGCAAVAAALLLYLLGARAQGGTRSPRCDCAGDEHKKIG
LFCCRGCPAGHYLKAPCTEPCGNSTCLVCPQDTFLAWENHHNSECARCQACDEQASQV
ALENCSAVADTRGGCKPGWFVECQVSQCVSSSFFYCQPCLDCGALHRHTRLLGSRRDT
DCGTCLPGFYEHGDGCVSCPTSTLGSCPERCAAVCGWRQSRWCAGNASGRTGMDRGEA
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/function="mediates apoptosis"
/note="LARD-2; similar to Fas (
soluble; truncated before the
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                                                                                                             Young, P.E., Augustus, M., Carter, K.C., Ebne Horrigan, S., Soppet, D.R. and Weaver, Z. Cancer gene determination and therapeutic
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Cancer gene
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Pharmaceuticals (US)
Location/Qualifiers
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                                         /organism="Homo sapiens"
/db_xref="taxon:9606"
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                                                          CATCTGTCACCCTTGGACAGCGCCCACACCCTTCTAGCACCTCCTGACAGCAGTGAGAAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ritson, J.
Direct Submission
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Best Local Similarity:
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Mammalia; Eutheria; Pr
1 (bases 1 to 1669)
Warzocha,K., Ribeiro, I
Salles,G.
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                       Metazoa; Chordata;
Eutheria; Primates;
          Ribeiro, P.,
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         Charlot, C.,
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3 beta
                      Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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(DR3)
         Renard, N.,
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         Coiffier, B.
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CysGlnValSerGlnCysValSerSerSerProPheTyrCysGlnProCysLeuAspCys
                                             HisasnSerGluCysalaargCysGlnalaCysaspGluGlnalaSerGlnValalaLeu
                                                                                                                                                                                        CysGlyAsnSerThrCysLeuValCysProGlnAspThrPheLeuAlaTrpGluAsnHis
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/product="death receptor 3 beta"
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/db_xref="gi:2570831"
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ALENCSAVADTRCGCKCGWFVECQVSQCYSSSPFYCQPCLDCGALHRHTKLLCSRRDT
DCGTCLPGYYEHGDGCVSCPTEPPSLAGAPWGAVQSAVPLSVAGGRVGVFWYQVLLAG
LVVPLLLGATLTYTYRHCWPHKPLVTADEAGMEALTPPPATHLSPLDSAHTLLAPPDS
SEKICTVQLVGNSWTPGYPETQBALCPQVTWSWPDQLPSRALGPARAPTLSPESPAGSP
AMMIQPGOLYDVMDAVPARRWKEFVRTLGLREAEIEAVEVEIGLFRDQQYEMLKHWR
QQQPAGLGAVYAALERWGLDGCVEDLRSRLQRGP"
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patients with non-Hodgkin's lymph
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                                            aAlaLeuGluArgMetGlyLeuAspGlyCysValGluAspLeuArgSerArgLeuGlnAr
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Result No.

SUMMARIES

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Pred. No. is the number of results predicted by chance to have score greater than or equal to the score of the result being predicted by analysis of the total score distribution. printed,

Result	No.	Score	Query Match	Length	DB	ID	ript
	ъ:	2209	100.0	1254	18	AAT89427	Death domain conta
	ນ	2209		1254	20	AAX00925	domain
	٠ ψ	2209		1254	21	AAC68777	
	<u>ء</u> س	2209	100.0	1634	22	AAT91180 AAH27782	Human genomic DNA
	6	2209		1634	24	AAL47186	
	7	2209		1662	22	AAC91477	
	0 00	2209	100.0	1662	24	ABK40265	encoding
	10	2209	100.0	1783	20 6	AAT89426	
	= ;	2209		1783	21	AAC68776	Human death domain
	12	2209	100.0	1847	19	AAV28700	
	13	2140.5	96.9	1743	24	ABL64119	Breast cancer rela
	л <u>н</u>	2094	94.8	1250	22	AAF83770	Nucleotide sequenc
	16	1498.5	67.8	10797	23	ABK42690	Genomic seguence #
	17	1373	62.2	1251	19	AAV28701	
	18	957.5	 	816	24	ABL67258	Thyroid cancer rel
	20	937	42.4	787	24	AAI,47187	Human rheumatoid a
	21	383.5	17.4	2130	24	ABK63694	equence difi
	2 2	3 6 9	16.7	1368	14	AAQ49932	Lambda-derived TNF
	24	369	16.7	2062	13	AA020973	TNF-alpha binding
	25	369	16.7	2062	13	AAQ24440	Encodes TNF-alpha
	26	369	16.7	2088	12	AAQ10883	TNF inhibito
	2 6	26 P	16.7	2088	2 2	AAC83946	Human 30 kDa TNF i
	29	369	16.7	2111	22	AAH48859	Human TNFBP-associ
	30	369	٠	2111	24	ABK84039	n cDNA d
	ب ا	3 6 6	16.7	2111	24	ABN95862	#2360 us
	ωü	369	16.7	2161	21	AAZ48475	Human tumour necro
	34	369	•	2161	24	ABK13194	an tumour
	u U	369	٠	17	16	AAQ90513	N.
	ט קר	167 S	16.7	2173	1 4	AAQ306784	
	38		o :	11	12	AAQ10955	Encodes human 55kD
	39	365		2176	12	AAQ12215	
,	40	320	14.5	11	24	ABQ47530	Oligonucleotide fo
a	<u> </u>	320	14.5	511	24	ABQ47531	Oligonucleotide fo
,	<u>ء</u> ۾	25.0	11.0	511 511	2 1	AAQU6282	- 2
•	4	253	11.5	511	24	ABO47533	Oligonucleotide fo
	45	229	10.4	1704	19	AAV34509	~

## ALIGNMENTS

RESULT 1
AAT89427
ID AAT8

AAT89427 standard; cDNA; 1254 BP.

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sig_peptide	Кеу	•	Homo sapiens.		inflammation;	Death domain		Death domain		02-MAR-1998		AAT89427;
172 /*tag= a	Location/Qualifiers				inflammation; NF-kappaB; ds.	Death domain containing receptor; DR3; human; apoptosis;	•	Death domain containing receptor DR3 cDNA.		02-MAR-1998 (first entry)		

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                                                                                                                                                                                                                                                                                                                    Score:
                                                                                                                                                                                                                                                                                                                            Alignment Scores: Pred. No.:
                                                                                                                                                                                                                                                                                                                                                                          This cDNA clone codes for human death domain containing receptor (C) DR3 (see AAW31517), a novel member of the tumour necrosis factor receptor family. It was isolated from a HUVEC cDNA library. Related death domain containing receptor DR3-VI cDNA (see AAT89426) (C) was isolated from a human testis tumour cDNA library. The genes isolated from a human testis tumour cDNA library. The genes (C) have also been identified in cDNA libraries of foetal liver, foetal brain, tonsil and leukocyte. Nucleic acids encoding full-length or mature DR3, or the extracellular, transmembrane, intraclelular or especially the death domain of DR3, can be used to produce recombinant polypeptides in transformed host cells. These polypeptides can be used to treat diseases and disorders associated with the inhibition of apoptosis. Antagonistscan be used to treat diseases and disorders associated with increased apoptosis and for treating inflammatory diseases and disorders.
                                                                                                                                                                                                                                                                                                                                                            Sequence 1254
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CysGlnValSerGlnCysValSerSerSerProPheTyrCysGlnProCysLeuAspCys
                      CTGTTTTGTTGCAGAGGCTGCCCAGCGGGGCACTACCTGAAGGCCCCTTGCACGGAGCCC
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DB; AAW31517.
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12-MAR-1996;
17-OCT-1996;
                                                                                                                                                                                                                                                    The invention provides nucleotide sequences encoding death domain containing receptor polypeptides DR3-V1 and DR3. The DR3-V1 cDNA clone is contained in ATCC deposition No. 97456 and the DR3 cDNA clone is contained in ATCC deposition No. 97757. Recombinant vectors comprising the nucleic acid sequences and optionally the leader sequences are used for the recombinant production of the proteins. The present sequence represents a cDNA encoding the death domain containing receptor polypeptide (DR3).
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                                                                                                                                                                                                                                                                                                                      Claim
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Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention provides the protein and coding sequences for two death domain containing receptors, designated DR3 and DR3 VI. These receptors are involved in apoptosis, and the sequences given can be used in the treatment of cancers, infections, cardiovascular disorders such as arrhythmias, ischaemia, aneurysms, arterial occlusive diseases, embolisms and congenital heart defects, neurodegenerative diseases including and congenital heart diseases, autoimmune disease such as multiple sclerosis, arthritis, diabetes, Graves' disease, asthma and psoriasis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1254
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Treating graft-versus-host disease, cancer, immunodeficiency or autoimmune disease comprising administering an antibody to Death Containing Receptor proteins and a second therapeutic agent -
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Location/Qualifiers

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                                                                                                                                                                                                                                                                                          cDNA clone FH20.57 (ATCC 55820) codes for a novel polypeptide (see W26709), designated Apo-3, that stimulates or induces apoptotic activity in mammalian cells. It was isolated from a human foetal heart cDNA library by screening with probes (see T91183-84) based on an EST sequence (GenBank locus W71984) that showed homology to the intracellular domain of human TMFR1 and CD95. Amino acid residues 1-181 of Apo-3 are identical to another novel apoptosis polypeptide, Apo-2LI (see W25708). Nucleic acids encoding Apo-3 can be used diagnostically for tissue-specific typing and to produce recombinant Apo-3 polypeptides, especially the extracellular domain (amino acids 1-198) or death domain (amino acids 1-199) or death domain (amino acids 1-199) or death domain (amino acids 1-199) or death domain containing cells that purposes. Non-human transgenic animals containing cells that express Apo-3 nucleic acid, and knockout animals containing cells that cells that have an altered Apo-3 gene, can be used in drug screening and development.
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                                                                                                                                                                                                                                                                                                represents DNA the diam
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                                                                                                                                                                                                                                                                                                                                                                         Diagnosing rheumatoid arthritis or comparing expression of mRNA
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27-MAR-2001;
30-MAR-2001;
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                                            GluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGlyTrpPheValGlu
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polypeptides. The PRO polypeptides, anti-PRO actus encouning PRO CC antagonists are useful for treating and diagnosing immune related CC disorders such as systemic lupus erythematosus, rheumatoid arthritis, CC systemic sterosis, idiopathic inflammatory myopathies, Sjogren's CC systemic vasculitis, sarcoidosis, autoimmune haemolytic canaemia, autoimmune thrombocytopaenia, thyroiditis, diabetes mellitus, CC immune-mediated renal disease, demyelinating diseases of the central cand peripheral nervous systems (such as multiple sclerosis, idiopathic inflammatory demyelinating polyneuropathy or Guillain-Barre syndrome, and chronic demyelinating polyneuropathy or Guillain-Barre syndrome, and chronic (such as infectious, autoimmune chronic ective hepatitis, primary cirrhosis, granulomatous hepatitis and sclerosing cholangitis), CC disease, autoimmune or immune-mediated skin diseases (such as bullous skin diseases, erythema multiforme, contact dermatitis, psoriasis), considered, such as asthma, allergic rhinitis, atopic dermatitis, food hypersensitivity and urticaria), immunological diseases of the cund diseases such as asthma, allergic rhinitis, atopic dermatitis, cand hypersensitivity pneumonitis), transplantation associated diseases including graft rejection and graft-versus-host diseases.
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17-MAY-2000;
22-MAY-2000;
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28-JUL-1999
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                                                                                                                                                                                                                                                                                                                                                               The
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                                                                                                                                                                                                                                                                                                                                                                                           Claim
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DB; AAB50918.
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2000WO-USO0376
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2000WO-USO04341
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2000WO-USO0414
2000WO-USO0884
2000WO-USO0377
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99US-0170262.
99WO-US30911.
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99WO-US28313.
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                                             polypeptides and the polynucleotide sequences encoding them. The PRO polypeptides, agonists, antagonists or anti-PRO antibodies are useful for treating benign or malignant tumours (e.g. renal, kidney, bladder, breast, etc), leukaemias and lymphoid malignancies, other disorders such as neuronal, glial, astrocytal, hypothalamic, glandul macrophagal, stromal and blastocoelic disorders, inflammatory, immun and angiogenic disorders. The polynucleotide sequences are also useful in gene therapy. ABK40254-ABK40288 encode for the human PRO polypeptides of the invention.
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                              Death domain containing receptor polypeptide(s) DR3 and DR3-V1 for activation of apoptosis and NF-kappaB, antagonists can be to treat inflammatory diseases
                                                                                                                                                                             Dillon PJ,
                                                                                                                     P-PSDB;
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                                                       US-09-993-234-6_COPY_25_417 (1-393) x AAT89426 (1-1783)
                                                                                                    Percent Similarity:
Best Local Similarity:
Query Match:
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                                                                                                                                               Score:
                                                                                                                                                            Pred.
                                                                                                                                                                        Alignment Scores:
                                                                                                                                                                                                                                Commain containing receptor DR3-VI (see AAW31516), a novel member of the tumour necrosis factor receptor family. It was isolated from a CDNA library derived from cells of a human testis tumour. Related feath domain containing receptor DR3 cDNA (see AAT89427) was isolated from a HUVEC cDNA library. The genes have also been identified in CDNA libraries of foetal liver, foetal brain, tonsil and leukocyte. Nucleic acids encoding full-length or mature DR3-VI can be used to produce recombinant polypeptides in transformed host cells. These polypeptides can be used to treat diseases and disorders associated with increased apoptosis. Antagonists, such as antibodies associated with increased apoptosis and for treating inflammatory
                                                                                                                                                           No.:
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and disorders.
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{\tt ThrProProProAlaThrHisLeuSerProLeuAspSerAlaHisThrLeuLeuAlaPro}
                                           CGCCACTGCTGGCCTCACAAGCCCCTGGTTACTGCAGATGAAGCTGGGATGGAGGCTCTG
                       ArgHisCysTrpProHisLysProLeuValThrAlaAspGluAlaGlyMetGluAlaLeu
                                                                                                                                                                                                                                                   GluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGlyTrpPheValGlu
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                                                                                                                                                              TGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGCCAACCATGCCTAGACTGC
                                                                                                                                                                      CysGlnValSerGlnCysValSerSerSerProPheTyrCysGlnProCysLeuAspCys
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Claim

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Page 71-73;

108pp; English

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ID AAXOO
XX AAXOO
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XX Death
XX Death
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P-PSDB; AAW95537.
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96US-0013285.
96US-0028711.
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SXCCCCCCCXPXPTX
                                                                               New death domain containing receptor and optionally comprising leader sequence
                                                                    Claim 2;
                                                                   Fig 1,
                                                                  50pp; Japanese.
                                                                                      recombinant vector
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The invention provides nucleotide sequences encoding death domain containing receptor polypeptides DR3-V1 and DR3. The DR3-V1 cDNA clone is contained in ATCC deposition No. 97456 and the DR3 cDNA clone is contained in ATCC deposition No. 97757. Recombinant vectors comprising the nucleic acid sequences and optionally the leader sequences are used for the recombinant production of the proteins. The present sequence represents a cDNA encoding the death domain containing receptor polypeptide (DR3-V1).

Sequence 1783 B₽; 330 A; 562 Ç 564 <u>ი</u> 327 7 0 other;

Alignment Pred. No.:

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LeuPheCysCysArgGlyCysProAlaGlyHisTyrLeuLysAlaProCysThrGluPro
                                                               GlySerCysProGluArgCysAlaAlaValCysGlyTrpArgGlnMetPheTrpValGln
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                                   ValLeuLeuAlaGlyLeuValValProLeuLeuGlyAlaThrLeuThrTyr
                                                       GGGAGCTGTCCAGAGCGCTGTGCCGCTGTCTGTGGCTGGAGGCAGATGTTCTGGGTCCAG
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ThrProProProAlaThrHisLeuSerProLeuAspSerAlaHisThrLeuLeuAlaPro

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RESULT 11
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(YUGG/)
    Treating graft-versus-host disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; death domain containing receptor; DR3-V1; cancer; autoimmune disorder; inflammation; cardiovascular disorder; infection;
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28-MAY-1999;
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autoimmune disease comprising administering an antibody to Death Domain Containing Receptor proteins and a second therapeutic agent -  $\,$ 

Example 1; Fig 1; 273pp; English.

The present invention provides the protein and coding sequences for two death domain containing receptors, designated DR3 and DR3-V1. These receptors are involved in apoptosis, and the sequences given can be used in the treatment of cancers, infections, cardiovascular disorders such as arrhythmias, ischaemia, aneurysms, arterial occlusive diseases, embolisms and congenital heart defects, neurodegenerative diseases including Alzheimer's and Parkinson's diseases, autoimmune disease such as multiple sclerosis, arthritis, diabetes, Grave', disease, asthma and psoriasis, promote angiogenesis and wound healing. embolisms multiple 25

Sequence 1783 BP; 330 A; 562 C; 564 G; 327 T; 0 other;

US-09-993-234-6\_COPY\_25\_417 (1-393) x AAC68776 (1-1783)

Query Match:

Percent Similarity: Best Local Similarity:

3.21e-117 2209.00 100.00% 100.00% 100.00% 100.00%

Conservative: Mismatches: Indels:

1783 393 0 0 0

Length: Matches:

Alignment Scores:

No.:

Qy 201 ArgHisCysTrpProHisLysProLeuValThrAlaAspGluAlaGlyMetGluAlaLeu	Qy 181 ValLeuLeuAlaGlyLeuValValProLeuLeuLeuGlyAlaThrLeuThrTyrThrTyr	Qy 161 GlySerCysProGluArgCysAlaAlaValCysGlyTrpArgGlnMetPheTrpValGln	QY 141 CysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysProThrSerThrLeu 	Qy 121 GlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThrAspCysGlyThr 	QY 101 CysGlnValSerGlnCysValSerSerFroPheTyrCysGlnProCysLeuAspCys	OY 81 GluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGlyTrpPheValGlu 	Qy 61 HisAsnSerGluCysAlaArgCysGlnAlaCysAspGluGlnAlaSerGlnValAlaLeu 	QY 41 CysG1yAsnSerThrCysLeuValCysProGlnAspThrPheLeuAlaTrpGluAsnHis	QY 21 LeuPheCysCysArgGlyCysProAlaGlyHisTyrLeuLysAlaProCysThrGluPro	Qy 1 GlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHlsLysLysIleGly
aAspGluAlaGlyMetGluAlaLeu 220	euGlyAlaThrLeuThrTyrThrTyr 200 	lyTrpargGlnMetPheTrpValGln 180 	ysValSerCysProThrSerThrLeu 160 	erArgArgAspThrAspCysGlyThr 140             CCGCAGAGATACTGACTGTGGGACC 722	heTyrCysGlnProCysLeuAspCys 120 	lyCysLysProGlyTrpPheValGlu 100 	spGluGlnAlaSerGlnValAlaLeu 80 	SPThrPheLeuAlaTrpGluAsnHis 60 	yrLeuLysAlaProCysThrGluPro 40 	laGlyAspPheHisLySLySIleGly 20               CGGTGACTTCCACAAGAAGATTGGT 362

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      DNA encoding apoptosis inducing receptor - which is transmembrane protein, useful for regulating cell de
                            P-PSDB;
                                                                                                                                                                                     Apoptosis inducing receptor; AIR protein; Type I transmembrane protein; tumour cell
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DB; AAW57045.
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death; autoimmune disease;
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Page 28-30; 45pp; English.

This sequence encodes the human apoptosis inducing receptor (AIR) of the invention. AIR is a Type I transmembrane protein, soluble forms of which can be used to regulate cell death in a therapeutic setting. Soluble AIR can also be used in vitro to block apoptosis or AIR-expressing cells, or to screen agonists or antagonists of AIR activity. The cytoplasmic domain of AIR can be used to develop assays for inhibitors of AIR-induced cell death, which is useful to regulate cell death in a therapeutic setting as well as in vitro. Agonists of AIR activity can be used to kill tumour cells that express AIR, or T cells expressing AIR in autoimmune diseases.

BP; 316 A; 605 C; 579 G; 347 T; 0 other;

3.33e-117 2209.00 100.00% 100.00% 100.00% 100.00%

Length:
Matches:
Conservative:
Mismatches:
Indels:

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US-09-993-234-6_COPY_25_417 (1-393)
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                                                                                                                         GTGCTCCTGGCTGGCCTTGTGGTCCCCCCTCCTGCTTGGGGCCACCCTGACCTACACATAC
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ProAspSerSerGluLysIleCysThrValGlnLeuValGlyAsnSerTrpThrProGly
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RESULT 13
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03-OCT-2000;
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28-SEP-2000;
29-SEP-2000;
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29-SEP-2000;
02-OCT-2000;
02-OCT-2000;
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2000US-237278P
2000US-237278P
2000US-237278P
2000US-237278P
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Soppet Young PE, Augustus M, Weaver Z; Carter KC, Ebner 'n Endress ્ Horrigan s

WPI; 2002-188264/24.

Screening for anti-neoplastic agent involves exposing cells to a chemical agent to be tested for anti-neoplastic activity, and determining a change in expression of a gene of a signature gene set

Claim 1; SEQ ID 2456; 44pp; English.

The present invention describes a method (M1) for screening for an CC anti-neoplastic agent. The method involves exposing cells to a chemical agent to be tested for anti-neoplastic activity, determining a change in CC expression of at least one gene (I) of a signature gene set, where (I) CC comprises a sequence (S) selected from 8447 sequences (given in ABIJ6164 CC to ABIJ7010), or is at least 95% identical to (S), where a change in CC expression is indicative of anti-neoplastic activity. (I) has cytostatic CC activity and can be used in gene therapy. M1 can be used for screening CC is the data collected with respect to the anti-neoplastic agent as a CC result of M1, and the data is sufficient to convey the chemical CC tructure and/or properties of the agent. M1 can be used in the CC treatment of cancer such as colon, breast, stomach, lung, thyroid, CC oesophageal, ovarian, kidney, prostate or pancreatic cancer, CC infiltrating lobular cancer, squamous cell carcinoma, neuroendocrine CC carcinoma, papillary carcinoma and Wilm's tumour.

Sequence 1743 BP; 347 A; 543 ç; 549 <u>.</u> 304 Ŧ; 0

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Query Match:
DB:
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2.5e-113
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      Length:
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Mismatches:
Indels:
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US-09-993-234-6\_COPY\_25\_417 (1-393)× ABL64119 (1-1743)

- γ В 141 20
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                                 GTGCGCACGCTGGGGCTGCGCGAGGCCAGAGATCGAAGCCGTGGAGGTGGAGATCGGTCTC
                                          ValArgThrLeuGlyLeuArgGluAlaGluIleGluAlaValGluValGluIleGlyArg
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The invention relates to a composition comprising a biologically active TR3-specific binding agent (I) that binds to TR3 and inhibits the proliferation of cells expressing TR3. (I) identified by the methods are useful for treating a subject suspected of having a disease associated with a proliferation of cells expressing TR3 especially leukemias or

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XX TR3;
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Alignment Scores:
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Best Local Similarity:
Query Match:
DB:
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                 789
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                       ProAspSerSerGluLysIleCysThrValGlnLeuValGlyAsnSerTrpThrProGly
                                                                                                                                                                 CCTGACAGCAGTGAGAAGATCTGCACCGTCCAGTTGGTGGGTAACAGCTGGACCCCTGGC
                                                  ACCCCACCACCGGCCACCCATCTGTCACCCTTGGACAGCGCCCACACCCCTTCTAGCACCT
                                                           ThrProProProAlaThrHisLeuSerProLeuAspSerAlaHisThrLeuLeuAlaPro
                                                                                                 ArgHisCysTrpProHisLysProLeuValThrAlaAspGluAlaGlyMetGluAlaLeu
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Alignment
Pred. No.:
Score:
US-09-993-234-6_COPY_25_417 (1-393)
                  Percent Similarity:
Best Local Similarity:
Query Match:
DB:
                                                                                      24-OCT-2000;
27-MAR-2001;
30-MAR-2001;
                                                                                                    The present invention relates to the human DR3 gene, which is associated with rheumatoid arthritis. Certain mutations in the gene can be linked to the disease. The sequences can be used to evaluate disease onset and its possibility and to provide therapy and remedies. The present sequence is the gene of the invention.
                                                                                                                                                                        Genomes, particularly DR3 genomic DNA, participating arthritis via mutation, useful in evaluating disease possibility and providing therapy and remedies -
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        CTCCCTGCCTGACCCACTCCTGTCCCCATGGTGACGCATGCCTCTCCTGGATTGCAGGTTC 1746
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                                         CCACACACGGCTACTCTGTGA-GTACCCCCACCCAGGGCTCTCTACTCCCAGACCCCCTT
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	2765	CCTCCCAAAGTGCTGGGATTACAGGCATGAGCCACCGCGCCCGGCCTCCATTCAAGTCTT	2706	Ъ
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	2705	GTTTCACCGTGTTAGCCAAGATGGTCTTGATCACCTGACCTCGTGATCCACCCCGCCTTGG	2646	DЬ
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	2645	GGGACTACAGGAGCCCACCACCCCCGGCTAATTTTTTGTATTTTAGTAGAGACGGG	2586	DЬ
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-	2585	S CACTACAACCTCTGCCTGGGTTCAAGTGATTCTCCTGGCTCAGCCTCCCAAGTAGCT	2526	В
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	2525	TTTTTGAGATGGAGACTTGCTCTGTCACCCAGGCTGGAGTGCAGTGGCACGATCTTGGCT	2466	Дb
	158		158	Qy
	2465	5 GATCAGAGGTAAGGGACTTTCTGTAGCTATTCAAGTCTTTTTTTT	2406	Db
	1:58		158	Qy
	2405	5 CTCTGATGTGTGGTGTCCCCTTTCTCAACATGGGATGTTAGCAGCTAATTCACAGGCCTTT	2346	В
	158		15	Qy
	2345	6 GAGTCAGGACTGTGCTAGTTCTCTAGGGGGATCTTGGGCAAGTGCAGAGAATTCGCGT	228	Дb
	158	8	15	Qy
	2285	6 TATTAAATCTATTAAATATTAAATATTAAATCTATTAAATAAATAAATAAATACAAAGGGCTGA	2226	фa
	158	8	15	Qγ
	2225	6 ATCGTGCCACTGCACTTCAGCCTGGGCAACAGAGCGAGACCTATTAAATAAA	216	ДЬ
	158	80	15	Qy
	2165	6 GTCACATTGATATAGCTTTAAAACTTGGGCTGAAGGAGGTTGAGGCTGCAGTGAGCTATG	2106	Дb
:	158	8	15	Qy
	2105	6 CAGCATTCCCCCCCAGCGATCCTAGCCAGATATTACAGATGATTTGTCACTTACACAGAGA	2046	DЬ
	158	8	15	Qy
	2045	6 AAAGAGCAATCTGGATCCGCCTTAGCCAGATACATAAGGGTATTTGCCTTCACTTTCAGC	1986	망
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	1985	6 GGAGGAGGGAGCCTTGGCCCTGTGATGGGTGGGCCCCACTTCAGGCAAACTTAGATGGCA	1926	Db
	158	8	158	Qy
	1925	6 CAGAGCAGGGGCCTGGGGTGGGGCAGGTGCTGCTGGTTCAGGAATAGGAAGAGGGGATAG	1866	ΔQ
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	158		152	Qy
	1806	7 CCGCAGAGATACTGACTGTGGGACCTGCCTGCCTGGCTTCTATGAACATGGCGATGGCTG	1747	DЪ
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3784	3725 CTGCCCTCAAGGGGTGCTCAGTCAGCTGGAGTTCAGATTCGTACACAGGAGCTAACAGTT 3	дb	
223	223 2	Qy	
3724	3665 AGCCCTAGTGGACAGAGGTGTTGGGAGCAGAGTCATCAGTGGATGAGACCAGCACAGTGC 3	DЬ	
223	223 2	Qy	
3664	3605 GAGCTGGAAGATCAAGCCTTACTATGATCCCTGGAGCTTGGCACGCGGCCAGCACCGGGT 3	Db	-
223	223 2	Qy	
3604	CGGTAAGAACCTCACTGTGTGATTCTGGGCTGCCTTCTG	DЬ	
223	217 etGluAlaLeuThrProPro 2	Qy	
3544		Db	
217		Qγ	
3484	3425 GAGGCAGGGAGTGTGGGGTGCAGAGGAACCCTAGAGGAGCTGTACCAGCACCCAGGTCCA	Db	
211	211 2	Оy	
3424	3365 AGCCTGGGGTCAGGATGGGTAGCCCAGAGTCTACTCAACCCTGATACAGAAGGGGAAACT	Db	
211		Qy	
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3304	3245 CCTGGCTGGCCTTGTGGTCCCCCCTCCTGCTTGGGGCCACCCTGACCTACACATACCGCCA	ДЪ	
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3244	GARGGGGGCAACCACCAACACCACCACCACCAGCTGCTTTCAGTG-TTCTGGGTCCAACGACCTCCACCACCACCACCACCACCACCACCACCACCACCA	da D	
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3 1 4	TAGGTGGTGTGCTGGGAATGCGAGTGGGAGAACTGGGATGGACCGAGGGAGG		
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3005	2946 TTGGGTAGAGTGAGAGGCCTGGGGAGGGAAGGGTCTCTAGGATGGAGCAGATGCTGGGCA	Db	
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2945	2886 GAAGAAGCTCAGTGAATAGTGTCTGTGGTTGTTACCAGGGACACAATGACAGGAACATTC	Db	
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Search completed: April 6, 2003, 16:52:30 Job time: 436.927 secs

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Minimum DB seq
Maximum DB seq
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-DB-Published_Applications_Na -QFMT=fastap_SUFFIX=rnpb -MINMATCH=0.1
-LOOPCL=0 -LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=biosum62
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-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
-FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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is derived by analysis of the total score distribution.
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/cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
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US-10-081-280-9
US-10-112-793-9
US-10-112-193-11
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222 222 222 232 244 244 256 266 277 277 277 277 277 277 277 277 27	Sequence 1, Appli Sequence 1389, Ap Sequence 1377, Ap Sequence 1577, Ap Sequence 292, Appli Sequence 5, Appli Sequence 5, Appli Sequence 5, Appli Sequence 5, Appli Sequence 5, Appli

## ALIGNMENTS

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RESULT 1
US-09-333-966-3
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 3, Apprince 1 Patent No. US20020009773A1 Patent NFORMATION:
                                                                                                                                                                                                                                                                        APPLICANT: Gentz, KEILL, APPLICANT: Dillon, Patrick J.
APPLICANT: Dillon, Patrick J.
TITLE OF INVENTION: Death Domain Containing Receptors
TITLE OF STATEMERS: 17
                      SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Yu, Guo-Liang APPLICANT: Ni, Jian APPLICANT: Dixit, Vishva
                                                                                                                                                                                             CITY: Washington
STATE: DC
                                                                                                                                                       COUNTRY: USA
ZIP: 20005-3934
                                                                                                                                                                                                                                      ADDRESSEE: Sterne, Kessler, Goldstein & Fox, P.L.L.C STREET: 1100 New York Ave., NW, Suite 600
  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Application US/09333966
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Gentz, Reiner L.
US/09/333,966
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Sequence

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US-09-993-234-6_COPY_25_417 (1-393) x US-09-333-966-3 (1-1254)
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Best Local Similarity:
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; LOCATION:
US-09-333-966-3
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INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 14:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
                                      493
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                                                               141
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FILING DATE: 12-MAR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Steffe, Eric K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 06-FEB-1997
PRIOR APPLICATION DATA:
            161
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APPLICATION NUMBER:
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STRANDEDNESS: doub
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                                               GlySerCysProGluArgCysAlaAlaValCysGlyTrpArgGlnMetPheTrpValGln
                                                                                                  GlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThrAspCysGlyThr 140
                               TGCCTGCCTGGCTTCTATGAACATGGCGATGGCTGCGTGTCCTGCCCCCACGAGCACCCTG
                                                                                                                                                     CysGlnValSerGlnCysValSerSerSerProPheTyrCysGlnProCysLeuAspCys
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                                                                                                                                                                                                                                                                                                  CysGlyAsnSerThrCysLeuValCysProGlnAspThrPheLeuAlaTrpGluAsnHis
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Steffe, Eric K.
Steffe, Bric K.
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Matches:
Conservative:
Mismatches:
Indels:
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Patent No. US20020165157A1
GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi
TITLE OF INVENTION: Apo-2
NUMBER OF SEQUENCES: 11
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                      COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 mb flops
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION NOMBER: US/10/081,280
FILING DATE: 21-Feb-2002
CLASSIFICATION: CUNknown>
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ADDRESSEE: Genente
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 APPLICATION DATA: APPLICATION NUMBER
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CITY: South San Francisco
STATE: California
                                                                                                                                                             COUNTRY: USA
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NUMBER: US/08/829,270
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INFORMATION FOR SEQ ID NO: 9:
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                                                                          GlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThrAspCysGlyThr 140
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                                        ArgHisCysTrpProHisLysProLeuValThrAlaAspGluAlaGlyMetGluAlaLeu
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TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
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REFERENCE/DOCKET NUMBER: P1007R1
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APPLICANT: Ashkenazi, Avi
TITLE OF INVENTION: Apo-2
NUMBER OF SEQUENCES: 28
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy di.
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION NUMBER: US/10/112,793
FILING DATE: 28-Mar-2002
CLASSIFICATION: <Unknown>
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                                                                         APPLICATION NUMBER: US/08/828,683A
FILING DATE: 31-Mar-1997
APPLICATION NUMBER: 08/625328
FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
ATTORNEY/AGENT INFORMATION:
             TELECOMMUNICATION INFORMATION TELEPHONE: 650/225-5416
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                       NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P10
COMMUNICATION: INFORMATION:
                                                                                                                                                                                                                                                                                                                         CITY: South San F
STATE: California
                                                                                                                                                                                                                                                                                                                COUNTRY: USA
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STREET: 1 DNA Way
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650/952-988:
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Best Local Similarity:
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TyrProGluThrGlnGluAlaLeuCysProGlnValThrTrpSerTrpAspGlnLeuPro
                        CGCCACTGCTGGCCTCACAAGCCCCCTGGTTACTGCAGATGAAGCTGGGATGGAGGCTCTG
                                                                                                     ArgHisCysTrpProHisLysProLeuValThrAlaAspGluAlaGlyMetGluAlaLeu
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TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
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Length:

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Alignment Scores: Pred. No.:
                                               ; SEQUENCE DESCRIPTION: SEQ US-10-112-193-11
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                                                                                                                                       INFORMATION FOR SEQ ID NO: 11: SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: 3.5 inch, 1.44 mb 1

COMPUTER: IEM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: WinPatin (Genentech)
                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
TELEFAX: 650/952-9881
                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Apo-3 POLYPEPTIDE NUMBER OF SEQUENCES: 15
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FILING DATE: 28-Mar-2002
CLASSIFICATION: <Unknown>
                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                   NAME: Marschang, Diane L. REGISTRATION NUMBER: 35,6
                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 08/928,069
FILING DATE: 11-Sep-1997
APPLICATION NUMBER: 60/026943
FILING DATE: 23-Sep-1996
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STREET: 1 DNA Way
                                                                                                                        LENGTH: 1634 base pairs
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                                                                                         Nucleic Acid
DEDNESS: Single
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ArgArgTrpLysGluPheValArgThrLeuGlyLeuArgGluAlaGluIleGluAlaVal
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Patent No. US20020146768A1
GENERAL INFORMATION:
APPLICANT: Ashkenazi,
                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 1634 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
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NUMBER OF SEQUENCES: 11
CORRESPONDENCE
                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/828,683
FILIUG DATE: <UNknown>
ATTORNEY/AGENT INFORMATION:
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ADDRESSEE: Genentech,
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                                                                                                                                                                                                                                                                                                  NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007P1
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FILING DATE: 19-No. US2002014676BA1-2001
CLASSIFICATION: <UNKNOWN>
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CITY: South San Francisco
STATE: California
                                                                                                                                                 TOPOLOGY: Linear NCE DESCRIPTION:
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TELEX: 910/371-7168
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Best Local Similarity:
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US-09-333-966-1
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                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 202-371-2540 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE:
APPLICATION NUMBER: NG
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APPLICANT: Yu, Gu
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                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 1783 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: 202-371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 60/013,285
FILLING DATE: 12-MAR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Steffe, Eric K.
REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 1488.031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 60/028,711 FILING DATE: 17-OCT-1996 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 06-FEB-PRIOR APPLICATION DATA:
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COMPUTER READABLE FORM:
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APPLICANT: Dixit, Vishva
APPLICANT: Gentz, Reiner L.
APPLICANT: Dillon, Patrick J.
TITLE OF INVENTION: Death Domain
NIMBER OF SEQUENCES: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                       NAME/KEY:
LOCATION:
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                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: both
1 GlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHisLysLysIleGly
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ZIP: 20005-3934
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CITY: Washington
STATE: DC
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1100 New York Ave.,
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202-371-2540
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NW, Suite
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Matches:
Conservative:
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FILE REFERENCE: 689290-77
CURRENT APPLICATION NUMBER: US/09/954,531
CURRENT APPLICATION NUMBER: US/60/233,133
PRIOR APPLICATION NUMBER: US/60/234,009
PRIOR APPLICATION NUMBER: US/60/234,009
PRIOR APPLICATION NUMBER: US/60/234,009
PRIOR APPLICATION NUMBER: US/60/234,034
PRIOR FILLING DATE: 2000-09-20
PRIOR FILLING DATE: 2000-09-20
PRIOR PRIOR APPLICATION NUMBER: US/60/234,509
PRIOR APPLICATION NUMBER: US/60/234,509
PRIOR APPLICATION NUMBER: US/60/234,567
PRIOR FILING DATE: 2000-09-2
SPENOR FILING DATE: 2000-09-2
SOFTWARE: PATENTAL NUMBER: US/60/234,567
PRIOR FILING DATE: 2000-09-2
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; TYPE: DNA
; ORGANISM: HOMO
US-09-954-531-1389
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GENERAL INFORMATION:
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                                     CysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysProThrSerThrLeu
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                                                                                               Sequence 1577, Application US/10092154 Publication No. US20030054375A1
 SEQ ID
                           APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PC009C1
CURRENT APPLICATION NUMBER: US/10/092,154
CURRENT FILING DATE: 2002-03-07
NUMBER OF SEQ ID NOS: 2003
Prior Application removed SOFTWARE: PatentIn Ver. 2. SEQ ID NO 1577
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          PatentIn Ver. 2.0

    See File Wrapper

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; TYPE: DNA
; ORGANISM: HOMO :
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aValAlaAspThrArgCysGlyCysLysProGlyTrpPheValGluCysGlnValSerGl
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244	7185 GGGACTACAGGAGCCCACCACCCCCGGCTAATTTTTTGTATTTTTAGTAGAGACGGG 7	DЬ
58	158 1	Qy
184	7125 CACTACAACCTCTGCTGCCTGGGTTCAAGTGATTCTCCTGCCTCAGCCTCCCAAGTAGCT 7	ఠ
58	158 1	Qy
124	7065 TTTTTGAGATGGAGACTTGCTCTGTCACCCAGGCTGGAGTGCAGTGGCACGATCTTGGCT 7	뮰
58	158 1	Qy
064	7005 GATCAGAGGTAAGGGACTTTCTGTAGCTATTCAAGTCTTTTTTTT	Db
58	158 1	Qy
004	6945 CTCTGATGTGTGGTGCCCTTTCTCAACATGGGATGTTAGCAGCCTAATTCACAGGCCTTT 7	Ъ
.58	158 1	Qy
944	6885 GAGTCAGGACTGTGCTGCTAGTTCTCTAGGGGATCTTGGGCAAGTGCAGAGAATTCGCGT 6	DЬ
58	158 1	Qy
884	6825 TATTAAATCTATTAAATATTAAATATTAAATCTATTAAATAAATAAATACAAAGGGCTGA 6	Db
58	158 1	Qy
824	6765 ATCGTGCCACTGCACTTCAGCCTGGGCAACAGAGCGAGACCTATTAAATAAA	В
58	158 1	Qy
764	6705 GTCACATTGATATAGCTTTAAAACTTGGGCTGAAGGAGGTTGAGGCTGCAGTGAGCTATG 6	뭥
58 `	158 1	Qy
704	6645 CAGCATTCCCCCCCAGCGATCCTAGCCAGATATTACAGATGATTTGTCACTTACACAGAGA 6	ф
58	158 1	Ş
644	6585 AAAGAGCAATCTGGATCCGCCTTAGCCAGATACATAAGGGTAITTGCCTTCACTTTCAGC 6	В
58	158 1	γo
584	6525 GGAGGAGGGAGCCTTGGCCCTGTGATGGGTGGGCCCCACTTCAGGCAAACTTAGATGGCA 6	Ъ
58	158	Qy
524	6465 CAGAGCAGGGGCCTGGGGTGGGGCAGGTGCTGCTTGAGGAATAGGAAGAGGGGATAG 6	Db
58	158 1	Qy
464		Ф
58	r 1	Qy
405	6346 CCGCAGAGATACTGACTGTGGGGACCTGCCTGCCTTCTATGAACATGGCGATGGCTG 6	В
52	ArgAspThrAspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCy 1	Qy
345	6286 CTCCCTGCCTGACCCACTCCTGTCCCATGGTGACGCATGCCTCTCCTGGATTGCAGGTTC 6	Ф
32	132 1	Qy
285	6227 CCACACACGGCTACTCTGTGA-GTACCCCCACCCAGGGCTCTCTACTCCCAGACCCCCTT 6	Db
32	HisThrArgLeuLeuCysSer 1	Qy
226	6167 ATGTGTCAGCAGTTCACCCTTCTACTGCCAACCATGCCTAGACTGCGGGGCCCTGCACCG 6	В
25	erSerProPheTyrCysGlnProCysLeuAspCysGlyAlaLeuHlsAr 1	Qy
166	6107 AGTGGCCGACACCCGCTGTGGCTGTAAGCCAGGCTGGTTTGTGGAGTGCCAGGTCAGCCA 6	В

8323	CCC IND I BUNCHUMUBI BI I BUBNUCNUNGI CAI CAB I BUNI BABNCCABCACAG TEC	Ň	- E	
	AGACCAGCAC AGTGC	8264	<u> </u>	
223		223	γQ	
8263	GAGCTGGAAGATCAAGCCTTACTATGATCCCTGGAGCTTGGCACGCGGCCAGCACCGGGT	8204	рь	<u></u>
223		223	Qy	
8203	TGGAGGCTCTGACCCCACCACCACCGGTAAGAACCTCACTGTGTGATTCTGGGCTGCCTTCTG	8144	Dр	
223	etGluAlaLeuThrProPro	217	Qy	
8143	GGAGGCTTGCCTGGTGGCTGACCGCAATCTCTCTGTGTCTGTC	8084	рь	
217		212	Qy	•
8083	GAGGCAGGGAGTGTGGGGTGCAGAGGAACCCTAGAGGAGCTGTACCAGCACCCAGGTCCA	8024	Db	
211		211	Qy	
8023	AGCCTGGGGTCAGGATGGGTAGCCCAGAGTCTACTCAACCCTGATACAGAAGGGGGAAACT	7964	В	
211		211	Qy	
7963	GCTGGCCTCACAAGCCCCTGGTTACTGGTAAGTACACACAC	7904	Db	
211	SCysTrpProHisLysProLeuValThr	202	Qy	
7903	CTGGCTGGCCTTGTGGTCCCCCTCCTGCTTGGGGCCACCCTGACCTACACATACCGCCA	7844	B . 8	
7843	3CAACCACCCAACACCCACCAGCTGCTTTCAGTG-TTCTGGGTCCAGGTGCT	o o	}	
182	GlnMetPheTrpValGlnVa	175	Qγ	
7784	TAGGTGGTGTGCTGGGAATGCGAGTGGGAGAACTGGGATGGACCGAGGGGAGGCGGGTGA	7725	рь	
174		174	Qy	
7724	GGCAGAG	7665	DЪ	
174	lyTrpArg	159	Qy	
7664	GTCTTAGGGAGCCCCTCCTGGCATGCACCCCCTCATCCCTCAGGCCACCCCCGTCCCTTG	7605	망	
158		158	Qγ	
7604	TTGGGTAGAGTGAGAGGCCTGGGGAGGGAAGGGTCTCTAGGATGGAGCAGATGCTGGGCA	7545	DЪ	
158		158	Qy	
7544	GAAGAAGCTCAGTGAATAGTGTCTGTGGTTGTTACCAGGGACACAATGACAGGAACATTC	7485	망	
158		158	Qγ	
7484	AAATAGGCAAAATCCCTGTCCTTTTGGGGTTGACATTCTAGTGACTCTTCATGTAGTCTA	7425	Дb	
158		158	γQ	
7424	TATTGAATATCTGCTATGTTCTACACACTGTTCTAGGTGCTGGGGGATGCAACAGGGGACA	7365	DЪ	
158		158	Qy	
7364	CCTCCCAAAGTGCTGGGATTACAGGCATGAGCCACCGCGCCCCGGCCTCCATTCAAGTCTT	7305	DЪ	
158		158	Qy	
7304	GTTTCACCGTGTTAGCCAAGATGGTCTTGATCACCTGACCTCGTGATCCACCCGCCTTGG	7245	ДD	
158		158	Qy	

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         LeuGlnArgGlyPro 393
CTGCAGCGCGGCCCG
                              GTTTACGCGGCCCTGGAGCGCATGGGGCTGGACGGCTGGGAAGACTTGCGCAGCCGC
                                       ValTyrAlaAlaLeuGluArgMetGlyLeuAspGlyCysValGluAspLeuArgSerArg
                                                                 AspGlnGlnTyrGluMetLeuLysArgTrpArgGlnGlnGlnDroAlaGlyLeuGlyAla
                                                                                                   CCGCAGCTCTACGACGTGATGGACGCGGTCCCAGCGCGGCGCTGGAAGGAGTTCGTGCGC
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RESULT 9

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SOFTWARE: Patentin Ver. 2
SEQ ID NO 1577
LENGTH: 10797
TYPE: DNA
ORGANISM: Homo sapiens
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APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteir
FILE REFERENCE: PC009
CURRENT APPLICATION NUMBER: US/09/764,847
CURRENT FILING DATE: 2001-01-17
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               CTTCCTGGTGACCTTCCCATCTCTCCATGTGCCCTTGGCCTCTGGTCGGCCTTAATCTCTG 5927
                                                                                                      CAGGCCTTTCTGAAGGAAGTGGCTGGCTCGGGCCCAAACTTGGGGTGTGAGGGTCCTGCA 5807
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                                                                                                                                                 GTGTCCCCAAGACACCTTCTTGGCCTGGGAGAACCACCATAATTCTGAATGTGCCCGCTG
                                                                                                                                                                                                             lCysProGlnAspThrPheLeuAlaTrpGluAsnHisHisAsnSerGluCysAlaArgCy 68
                                                                                                                                                                             sGlnAlaCysAspGlu-----
                                                                                                                                                                                                                                                                                        GCAGGCAGGGCTGGAGAGGTGGCGGGCAGGCCCGGGAGGTAAGAGGAGGCTGGCAGGGGA 5447
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	- 158		158	Qγ
	T 7004	CTCTGATGTGTGGTGTCCCTTTCTCAACATGGGATGTTAGCAGCTAATTCACAGGCCTTT	6945	Ъ
	- 158		158	Qγ
	T 6944	GAGTCAGGACTGTGCTGCTAGTTCTCTAGGGGATCTTGGGCAAGTGCAGAGAATTCGCG	6885	日
	- 158		158	Qy
	A 688,4	TATTAAATCTATTAAATATTAAATATTAAATCTATTAAATAAATAAATAAATACAAAGGGCTG	6825	В
	- 158		158	Ω
	A 6824	ATCGTGCCACTGCACTTCAGCCTGGGCAACAGAGCGAGACCTATTAAATAAA	6765	망
	- 158		158	δ
	G 6764	GTCACATTGATATAGCTTTAAAACTTGGGCTGAAGGAGGTTGAGGCTGCAGTGAGCTATG	6705	밁
	- 158		158	Qγ
	A 6704	CAGCATTCCCCCCAGCGATCCTAGCCAGATATTACAGATGATTTGTCACTTACACAGAG	6645	₽
	- 158		158	Ş
	C 6644	AAAGAGCAATCTGGATCCGCCTTAGCCAGATACATAAGGGTATTTGCCTTCACTTTCAG	6585	В
	- 158		158	δ
	A 6584	GGAGGAGGGAGCCTTGGCCCTGTGATGGGTGGGCCCCACTTCAGGCAAACTTAGATGGC	6525	ઠ
	- 158		158	Ş
	3 6524	CAGAGCAGGGGCCTGGGTGGGCAGGTGCTGGTTCAGGAATAGGAAGAGGGGGATAG	6465	В
	- 158		158	Q
	G 6464	GTGTCCTGCCCCACGTA-ATTCCTAGCTGTCGTGGGATGGAGGGAAGGGCGGCTGGGA	6406	Db
	- 158	SerCysProThrSer	152	δĀ
	G 6405	AGAGATACTGACTGTGGGACCTGCCTGCCTGGCTTCTATGAACATGGCGATGGCT	6346	망
	y 152	gAspThrAspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyC	133	Qγ
	3 6345	CTCCCTGCCTGACCCACTCCTGTCCCATGGTGACGCATGCCTCTCCTGGATTGCAGGTTC	6286	당
	- 132		132	QΥ
	T 6285	CCACACACGGCTACTCTGTGA-GTACCCCCACCCAGGGCTCTCTACTCCCAGACCCCCTT	6227	망
	- 132	ThrArgLeuLeuC	125	δ
	G 6226	ATGTGTCAGCAGTTCACCCTTCTACTGCCAACCATGCCTAGACTGCGGGGCCCTGCACCG	6167	밁
	r 125	SerSerProPheTyrCysGlnProCysLeuAspCysGlyAlaLeuHisA	105	VΩ
, į	1 105 A 6166	aValAlaAspThrArgCysGlyCysLysProGlyTrpPheValGluCysGlnValSerGJ 	85 6107	유 성
	C 6106	ACTCCCCATCTCCCTGCACCCCACCAGCC-TCCCAGGTGGCGCTGGAGAACTGTTCAG	6048	В
	85	lnAla	74	δ
	6047	ATTATCTTGGGATAATGCCTCTGCCTCTCCATGGGAGCCTTTGGCCCTGACTAACTCTCC	5988	Ъ
	73		73	δĀ
	5987	AGCTTCTCTCTTTTTAGGGTAGCCCTGTACCTGTCTGTCT	5928	용

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CAGGAGCACCCTGGGAATGCCAAATCTCTCTGTGAGAGCCTGTCAGAAGCACCCAGAAGCCCTGGGAACCCCTGGGAACCCCTGGGAACCCCTGGGAACCCCTGGGAACCCCTGGGAACCCCTAGAGCCCTTGTGCCGCTGTCCCAGAAGCCCTTGCAGAGCAACTAGAACTAGAACTAGAACTAGAACCTAGAACTAGAACTAGAACTAGAACTAGAACTAGAACTAGAACTAGAACTAGAACTAGAACTAGAACTAGAACTAGAACTAGAACTAGAACTACAACAACCAAC	GGGTAGAGTGAGAGGCCTGGGGAGGGAAGGGTCTCTAGGATGGAGGAGAGATGCTGGGCA	AAATAGGCAAAATCCCTGTCCTTTTGGGGTTGACATTCTAGTGACTCTTCATGTAGTCTA 748  58  69  60  60  60  60  60  60  60  60  60	TCCCAAAGTGCTGGGATTACAGGCATGAGCCACCGCGCCCGGCCCTCCATTCAAGTCTT 7	TGG	GATCAGAGGTAAGGGACTTTCTGTAGCTATTCAAGTCTTTTTTTT

	348 9223	ThrLeuGlyLeuArgGluAlaGluIleGluAlaValGluValGluIleGlyArgPheArg	329 9164	₽ 2
	328 9163	ProGlnLeuTyrAspValMetAspAlaValProAlaArgArgTrpLysGluPheValArg	309 9104	р У
	308 9103	AlaProThrLeuSerProGluSerProAlaGlySerProAlaMetMetLeuGlnProGly	289 9044	B 5
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	284		284	QΥ
	8863	AAGGGACATCAGTGGCCTGAGGCCTTGACCCCCATTCTCCTGTCTGCGGTGGGGAAGTTGTG	8804	Ъ
	284		284	Qy
	284 8803	GAGGCGCTCTGCCCGCAGGTGACATGTCCTGGGACCAGTTGCCCAGCAGAGCTCTTGGT	4- 0	B 2
	- 7	AAGATCTGCACCGTCCAGTTGGTGGGTAACAGCTGGACCCCTGGCTACCCCGAGACCC	366	g B
	265	Lys11eCysThrVa1G1nLeuVa1G1yAsnSerTrpThrProG1yTyrProG1uThrG1	24	Q.
	8683	CCATCTGTCACCCTTGGACAGCGCCCACACCCT	862,	рь
	245	ThrHisLeuSerProLeuAspSerAlaHisThrLeuLeuAlaProProAspSerSerC	226	Qy
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	225		224	Qγ
	8563	4 GTCATTGCGGGGGTGTCTGCTGACTGAACCAGGGACTGTCCCCTCCTGGAGAGGCACTGC	8504	Ъ
	223	3	22	Qγ
	8503	4 TCAAGGCAGAAGCAAGAGTTCTGCTGGGCTACAGTGAGAGCAGGGCCAACTGTGGGAGGT	844	Вþ
	223	3	22	Qy
	8443	4 CAATGGAAGGAGAGCCCCATGGTGCTGGGGGACAAGAGGAAGGA	838	В
	223	3	22	Qy
	8383	4 CTGCCCTCAAGGGGTGCTCAGTCAGCTGGAGTTCAGATTCGTACACAGGAGCTAACAGTT	832	DЬ
	223	3	22	Qγ
	8323	4 AGCCCTAGTGGACAGAGGTGTTGGGAGCAGAGTCATCAGTGGATGAGACCAGCACAGTGC	826	В
	223	3	22	Qy
	8263	4 GAGCTGGAAGATCAAGCCTTACTATGATCCCTGGAGCTTGGCACGCGCGCCAGCACCGGGT	820	밁
	223	3	22	Qγ
	82		814	뫄
	223	etGluAlaLeuThrPro	217	Qy

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; LENGTH: 816
; TYPE: DNA
; ORGANISM: Homo s
US-09-964-824A-292
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CURRENT FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: US/60/236,033
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR FILING DATE: 2000-09-28
UNMBER OF SEQ ID NOS: 583
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SEQ ID NO 292
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US-09-964-824A-292
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Cancer Gene Determination TITLE OF INVENTION: Cancer Gene Determination FILE OF INVENTION: Sets FILE REFERENCE: 689290-73
CURRENT APPROVALE-
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                                                             CysGlnValSerGlnCysValSerSerSerProPheTyrCysGlnProCysLeuAspCys 120
                                                                                                          GAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCCAGGCTGGTTTGTGGAG
                                                                                                                          GluAsnCysSerAlaŸalAlaAspThrArgCysGlyCysLysProGlyTrpPheValGlu 100
                                                                                                                                                                                        HisasnSerGluCysalaArgCysGlnAlaCysAspGluGlnAlaSerGlnValAlaLeu
                                                                                                                                                                      CATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGCCTCCCCAGGTGGCGCTG
                                                                                                                                                                                                                                                 CysG1yAsnSerThrCysLeuValCysProG1nAspThrPheLeuAlaTrpG1uAsnHis
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Conservative:
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; SEQUENCE DESCRIPTION: SEQ ID US-10-081-280-5
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US-10-081-280-5
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INFORMATION FOR SEQ ID NO: 5:
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                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/829,270
FILING DATE: 31-Mar-1997
ATTORNEY/AGENT INFORMATION:
NAME: MAISCHARD, Diane L.
REGISTRATION NUMBER: 35,600
REGISTRATION NUMBER: 35,600
                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/081,280
FILING DATE: 21-Feb-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: P1007R1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-5416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy
                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech,
                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Apo-2 LI AND Apo-3 POLYPEPTIDES NUMBER OF SEQUENCES: 11
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CITY: South San Francisco
STATE: California
COUNTRY: USA
ZIP: 94080
                                    TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech)
                                                                                        LENGTH: 1438 base pairs
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US-10-112-793-5
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GENERAL INFORMATION:
APPLICANT: AShkenazi, Avi J.
APPLICANT: MYENTION: Apo-2 LI AND
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          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/10/112,793
FILING DATE: 28-Mar-2002
CLASSIFICATION: <UNKnown>
PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA:
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MEDIUM TYPE: 3.5 1
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ADDRESSEE: Genentech,
STREET: 1 DNA Way
                                                                                                                                                                                                                                                 TITLE OF INVENTION: Apo-2
NUMBER OF SEQUENCES: 28
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CITY: South San Fi
STATE: California
APPLICATION NUMBER: US/08/828,683A
                                                                                                                                                     ZIP: 94080
                                                                                                                                                                    COUNTRY:
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Query Match:
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         Sequence 5, Application US/10112193 Publication No. US20030004313A1 GENERAL INFORMATION:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
TELEPHONE: 650/252-9881
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1438 base pairs
LENGTH: 1438 base pairs
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 APPLICANT:
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                                                                                               GlySerCysProGluArgCysAlaAlaValCysGlyTrpArgGln 175
                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
                                                                                                                                                                                            GlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThrAspCysGlyThr
                                                                                                                                                                                                                               TGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGCCAACCATGCCTAGACTGC
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STRANDEDNESS: Single
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 31-Mar-1997
APPLICATION NUMBER: 08/625328
FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
Ashkenazi, Avi J
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Matches:
Conservative:
Mismatches:
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US-09-993-234-6_COPY_25_417 (1-393) x US-10-112-193-5 (1-1438)
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Best Local Similarity:
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SEQUENCE CHARACTERISTICS:
LENGTH: 1438 base pairs
               101 CysGlnValSerGlnCysValSerSerSerProPheTyrCysGlnProCysLeuAspCys 120
                                                                              81 GluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGlyTrpPheValGlu 100
                                                                                                                            APPLICATION NUMBER: 08/928,069
FILING DATE: 11-Sep-1997
APPLICATION NUMBER: 60/026943
FILING DATE: 23-Sep-1996
ATTORNEY/AGENT INFORMATION:
NAME: MATSCHARG, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: 91052R1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                          LeuPheCysCysArgGlyCysProAlaGlyHisTyrLeuLysAlaProCysThrGluPro 40
TGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGCCAACCATGCCTAGACTGC
                                                              GAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGGCTGAAGCCAGGCTGGTTTGTGGAG
                                                                                                                                                                                                                                                           CTGTTTTGTTGCAGAGGCTGCCCAGCGGGGCACTACCTGAAGGCCCCTTGCACGGAGCCC
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STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: Linear
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STATE: California
COUNTRY: USA
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US-09-884-733-5
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Best Local Similarity:
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Patent No. US20020123116A1
GENERAL INFORMATION:
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               21 LeuPheCysCysArgGlyCysProAlaGlyHisTyrLeuLysAlaProCysThrGluPro
                                                                               1 GlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHisLysLysIleGly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/884,733
FILING DATE: 19-Jun-2001
CLASSIFICATION: -Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: 415/225-5416
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ADDRESSEE: Genentech,
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TITLE OF INVENTION: Apo-2 Ligand Inhibitor
NUMBER OF SEQUENCES: 9
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                                                              CAGGGCGCCACTCGTAGCCCCAGGTGTGACTGTGCCGGTGACTTCCACAAGAAGATTGGT 508
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                                                                                                                                                                                                                                                                                                                            TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
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APPLICATION NUMBER: 09/304,003
FILING DATE: 14-JUNE-2000
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CITY: South San Francisco
STATE: California
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US-09-993-234-5
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US-09-993-234-5
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                                                                                                                 TELEX: 910/371-71 INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech) CURRENT APPLICATION DATA:
                                                                                                                                                                    REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION: TELEPHONE: 415/225-5416
                                                                                                                                                                                                                                                       APPLICATION NUMBER: 08/828,683 FILING DATE: <Unknown> ATTORNEY/AGENT INFORMATION:
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              SEQUENCE DESCRIPTION:
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ADDRESSEE: Genentech,
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TITLE OF INVENTION: Apo-2
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                                                           LENGTH: 1438 base pairs
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FILING DATE: 19-No. US2002014676BA1-2001
CLASSIFICATION: <Unknown>
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CITY: South San Francisco
STATE: California
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                                              STRANDEDNESS: Single
                                                                                                                                                                                                                    NAME: Marschang, Diane L. REGISTRATION NUMBER: 35,600
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                                TOPOLOGY: Linear
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      161 GlySerCysProGluArgCysAlaAlaValCysGlyTrpArgGln 175
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                                             GlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThrAspCysGlyThr 140
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Search completed: April 7, 2003, 03:58:32 Job time: 188.643 secs

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Command line parameters:

-MODEL-frame+_p2n.model -DEV-xlh
-Q-/cgn2_1/USPTQ_spool/US09993234/runat_27032003_115456_15362/app_query.fasta_1.2346
-Q-/cgn2_1/USPTQ_spool/US09993234/runat_27032003_115456_15362/app_query.fasta_1.2346
-DP-EST -QEMT-fastap -SUFFIX-rst -MINMATCH=0.1 -LOOPCL=0 -LOOPEXT=0
-UNITS-bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS-human40.cd1 -LIST=45
-DOCALIGN=300 -THR_SCORE-pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15 -MODE-LOCAL
-OUTFMT-pto -NORM-ext -HEAPSIZE=500 -MILLEN=0 -MAXLEN=200000000
-USER=-US09993234_eCGN_1_1_4749_erunat_27032003_115456_15362 -NCPU=6 -ICPU=3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
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Listing first 45 summaries
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Maximum DB seq length: 200000000
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Perfect score:
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Ygapop 10.0 ,
Ygapop 6.0 ,
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Delop 6.0 ,
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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## ALIGNMENTS

REFERENCE AUTHORS TITLE		ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION		DEFINITION	Locus	BQ068309	RESULT 1
<pre>1 (bases 1 to 1010) NIH-MGC http://mgc.nci.nih.gov/. National Institutes of Health, Mammalian Gene Collection (MGC)</pre>	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.	Homo sapiens	human.	EST.	BQ068309.1 GI:19897355	BQ068309	5', mRNA sequence.	ᄛ	BQ068309 1010 bp mRNA linear EST 02-APR-2002		

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                                                                GATGGCTGCGTGTCCTGCCCCACGAGCACCCTGGGGAGCTGTCCAGAGCGCTGTGCCGCT
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Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at:
http://lmage.lnl.gov
Plate: LLAM12833 row: n column: 03
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                                                                                                                                                                                                                                                                                                                                                                                                                 Email: cgapbs-remail.nih.gov
This clone is available royalty-free through LLNL; contact the 'info@image.llnl.gov') for further information.
                                                                                                                                                                                                                                                                                                                                                                                                           IMAGE Consortium (info@image.llnl.gov)
Insert Length: 1568 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tumor Gene Index
Unpublished (1997)
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National Cancer Institute, Cancer Genome Ana
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wf52c03.x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone
IMAGE:3359204 3' similar to SW:WSL1_HUMAN Q93038 WSL-1 PROTEIN
PRECURSOR ;, mRNA sequence.
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1 (bases 1 to 698)
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/note="Organ: pooled; Vector: pT773D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Equal amounts of plasmid DNA from three normalized libraries (fetal lung MbHL19W, testis NHT, and B-cell NCI_CGAP_GCB1) were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 297480-302087, 682632-687239, 726408-728711, and 729996-731399. Subtraction by Bento Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                     /organism-"Homo sapiens"
/db_xref="taxon:9606"
/clone="Indage:2359204"
/clone_lib="Soares_NFL_T_GBC_S1"
/lab_host="DH10B"
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                                                                                                                                 GlySerCysProGluArgCysAlaAlaValCysGlyTrpArgGlnMetPheTrpValGln
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EST.
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Eukaryota; Metazoa;
Mammalia; Eutheria;
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  Chordata;
Primates;
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  Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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ThrLeuSerProGluSerProAlaGlySerProAlaMetMetLeuGlnProGlyProGln
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert Strausberg, Ph
Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 (bases 1 to 681)
NIH-MGC http://mgc.nci.nih.
National Institutes of Heal
Unpublished (1999)
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Plate: LLCM382 row: f column:
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/lab_host="DH10B (phage-resistant)"
/lab_host="DH10B (phage-resistant)"
/lab_host="Drgan: pancreas; Vector: pOTB7; Site_1: XhoI;
Site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the
following 5' adaptor: GGCACGAG(G). Library constructed
by Ling Hong in the laboratory of Gerald M. Rubin
(University of California, Berkeley) using ZAP-cDNA
synthesis kit (Stratagene) and Superscript II RT (Life
Technologies)."
Technologies."
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/clone_lib="NIH_MGC_39"
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/db_xref="taxon:9606"
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Health, Mammalian Gene
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Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLN
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC Clone distribution information
found through the I.M.A.G.E. Consortium/LLNL at:
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, M
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/clone="IMAGE:5214568"
/clone_lib="NIH_MGC_118"
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/lab_host="DH10B"
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        Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL;
IMAGE Consortium (info@image.llnl.gov) for further i
Possible reversed clone: similarity on wrong strand
Possible reversed clone: polyT not found
Seg primer: -40UP from Gibco
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Mammalia; Eutheria;
1 (bases 1 to 632)
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Contact: Robert Strausberg, Ph.D.
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US-09-993-234-6_COPY_25_417 (1-393) x BM009354 (1-651)
                                                                                                                                                                                                                                                     142 LeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysProThrSerThrLeuGly 161
                                                                                                                                                                                                                               4 CTGCCTGGCTTCTATGAACATGGCGATGGCTGCGTGTC-TGCCCCACGAGCACCCTGGGG
HisCysTrpProHisLysProLeuValThrAlaAspGluAlaGlyMetGluAlaLeuThr
                                                            Homo sapiens
Homo sapiens
Eukaryota; Metazoa; Chordata;
Eukaryota; Metazoa; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Robert Strausberg, Ph.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (1999)
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National Institutes of Health, M
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     late: LLCM1921 row: 1 column: lgh quality sequence stop: 649.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /tisue_type="amelanotic melanoma, cell line"
/tab_host="DH10B (phage-resistant)"
/note="Organ: skin; Vector: pOTB7; Site_1: XhoI; Site_2:
ECORI; CDNA made by oligo-dT priming. Directionally cloned
into EcoRI/XhoI sites using the following 5' adaptor:
GCCACGAG(G). Library constructed by Ling Hong in the
laboratory of Gerald M. Rubin (University of California,
BerKeley) using ZAP-CDNA synthesis kit (Stratagene) and
Superscript II RT (Life Technologies). Note: this is a
NIH_MGC Library."
a 224 c 204 g 116 t
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/db_xref="taxon:9606"
/clone="IMAGE:5443659"
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K-EST0076219 S22SNU16n1 H
5', mRNA sequence.
BM794760
BM794760.1 GI:19142992
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1 (bases 1 to 647)

Kim,N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim,M.R.
                                                                                                                                                                                                                                  Email: yongsung@mail.kribb.re.kr
Plate: 48 row: G column: 11
High quality sequence stop: 647.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                             Korea Research Institute of Bioscience 52 Eoeun-dong Yuseong-gu, Daejeon 305-3 Tel: +82-42-860-4470
                                                                                                                                                                                                                                                                                                                                                            Genome Research Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                              Contact: Kim YS
                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (2002)
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                                                                                                                                    /organism="Homo sapiens"
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/clone_lib="$22SNU16n1"
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                                                                          /cell_type="Lymphoblast-like"
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/lab_host="DH10B"
                                                                                                                          /tissue_type="Ascites"
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Park,H.S., Kim
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                              2.;, mRNA :
AI424936
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Mismatches:
Indels:
Gaps:
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Craniata;
Vertebrata;
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Euteleostomi;
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           CysGlnValSerGlnCysValSerSerSerProPheTyrCysGlnProCysLeuAspCys
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                                                                                                                    HisasnSerGluCysalaArgCysGlnAlaCysAspGluGlnAlaSerGlnValalaLeu
                                                                                                                                                       CATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGCCTCCCAGGTGGCGCTG
                                                  GAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCCAGGCTGGTTTGTGGAG
TGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGCCAACCATGCCTAGACTGC
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Unpublished (1997)
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   113
                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone_lib="NCI_CGAP_CLL1"
/tlssue_type="B-cell, chronic lymphotic leukemia"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:2109207"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   987a04.x1 Soares_fetal_heart_NbHH19W Homo sapiens cDNA clone IMAGE:1693710 3' similar to SW:WSL1_HUMAN Q93038 WSL-1 PROTEIN PRECURSOR ; mRNA segmence
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert Strausberg, Ph.D.

Email: cgapbs-r@mail.nih.gov

This clone is available royalty-free through LLNL; contact the

IMAGE Consortium (info@image.llnl.gov) for further information.

Insert Length: 1640 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                          Seq primer: -40m13 fwd. ET from Amersham
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Unpublished (1997)
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National Cancer Institute, Cancer Genome Anat
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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NbHL19W."
a 182 c
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/db_xref="taxon:9606"
/clone="IMAGE:1693710"
                                                                                                                                                                                                                                                                                                                          /dev_stage="19 weeks"
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                                                                                                                                                                                                                                                                                                Mammalia; Eutheria; Primates; Catarrhini; Homini (bases 1 to 539)
Wistow,G.J., Bernstein,S., Behal,A. and Smith,D. NEIBANK: EST analysis and bioinformatics for ocu Invest. Ophthalmol. Vis. Sci. 41, (2000) In pre Contact: Wistow G
                                                                                                                                                                                                                                                                                                                                                                                                                                                 BF726557 539 bp mRNA linear EST 05-JAN-2001 by08d05.y1 Human Lens cDNA (Un-normalized, unamplified): BY Homo sapiens cDNA clone by08d05 5', mRNA sequence.
BF726557
                                                                                                                                                                                                                                                            National Eye Institute 6/331, NIH, Bethesda, 1
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Eukaryota; Metazoa;
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                                                                                                                                                                                                        Email: graeme@helix.nih.gov
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primer: M13RP1 reverse primer
Location/Qualifiers
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301 496 0078
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BY"
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Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eukaryota; Metazoa; Chordata; Catarrhini; Hominidae; 1 (bases 1 to 552) 1 (bases 1, Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,
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                                                                                                                 mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the pCMVSPORT6 vector was constructed at Life Technologies , essentially following the protocols of the SuperScript Plasmid System full details of which are contained in the manufacturer's Instruction manual (http://www.lifetech.com/). First strand synthesis was carried out using a Not I primer-adapter [5'-pGACTAGTTCTAGATCGCGAGCGGCCC(T)15-3']. Not I/blunt end inserts were cloned into the Not I/EcoR V sites in the vector. EST analysis was performed on the unamplified library at the NIH Intramural Sequencing Center (NISC)." 97 a 192 c 167 g 83 t
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                   GluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysProGlyTrpPheValGlu 100
                                                                                                    HisAsnSerGluCysAlaArgCysGlnAlaCysAspGluGlnAlaSerGlnValAlaLeu 80
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GAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCCAGGCTGGTTTGTGGAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: yongsung@mail.kribb.re.kr
Plate: 31 row: H column: 06
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+82-42-860-4409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      bacterial alkaline phosphatase (BAP) and then decapped with tabacco acid pyrophosphatase (TAP). The decapped intact mRNA was ligated with DNA-RNA linker including EcoR I site by treatment of T4 RNA ligase and the first strand cDNA was synthesized from ollgo dT-selected mRNA by priming with dT-tailed vector. The dT-tailed vector was adjusted to have about 60nt. The cDNA vector was circularized with E. coli DNA ligase after digestion of EcoRI which site is also included in vector. An RNA strand converted to a DNA strand by Okayama-Berg method. The obtained cDNA vectors were used for transformation of competent cells E. coli ToplOF' by electroporation method. The cDNA libraries constructed by this method are full-length enriched cDNA libraries constructed by this method are 172 c 173 g 115 t
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Site_2: NotI; The poly (A)+ RNA was dephosphorylated with
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/db_xref="taxon:9606"
/clone="S6SNU620-31-H06"
/clone_lib="S6SNU620"
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/cell_line="SNU-620"
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                                                                                                                                                                                                                                                                                                                                                                                                                This clone is available royalty-free through LLNL; IMAGE Consortium (info@image.llnl.gov) for further i Possible reversed clone: similarity on wrong strand Possible reversed clone; polyT not found
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
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National Cancer Institute, Cancer Genome Ana
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/db_xref="taxon:9606"
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Kim, N.S., Hahn, Y.,
                                                                                                                                                                   Email: yongsung@mail.kribb.re.kr
Plate: 17 row: C column: 06
High quality sequence stop: 565.
Location/Qualifiers
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Contact: Kim YS
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/db_xref="taxon:9606"
/clone="S6SNU620-17-C06"
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/cell_line="SNU-620"
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Oh, K.J., Cheong, J.E., Sohn, H.Y., Kim, J.M., Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 514)
Kim, N.S., Hahn, Y.,
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Plate: 27 row: G column:
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l: +82-42-860-4470
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/db_xref="taxon:9606"
/clone="S6SNU620-27-G03"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Korea Research Institute of Bioscience & Biotechnology 52 Eoeun-dong Yuseong-gu, Daejeon 305-333, South Korea Tel: +82-42-860-4470
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kim, N.S., Hahn, x., ou,
Oh, K.J., Cheong, J.E.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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           /note="Organ: Stomach; Vector: pCNS; Site_1: EcoRI; Site_2: NoLI; The poly (A)+ RNA was dephosphorylated with bacterial alkaline phosphatase (BAP) and then decapped with tabacco acid pyrophosphatase (TAP). The decapped intact mRNA was ligated with DNA-RNA linker including EcoR I site by treatment of T4 RNA ligase and the first strand cDNA was synthesized from oligo dT-selected mRNA by priming with dT-tailed vector. The dT-tailed vector was adjusted to have about 60nt. The cDNA vector was circularized with E. coli DNA ligase after digestion of EcoRI which site is also included in vector. An RNA strand converted to a DNA strand by Okayama-Berg method. The obtained cDNA vectors were used for transformation of competent cells E. coli ToplOF' by electroporation method. The cDNA libraries constructed by this method are full-length enriched cDNA library."
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/db_xref="taxon:9606" .
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Search completed: April 6, 2003, 23:25:13 Job time : 2464.26 secs

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-LIST-45 -DOCALIGN-200 -THR_SCORE-pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15
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Encodes TNF-alpha
30kD TNF inhibitor
Human 30 kDa TNF i
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                                                                                                                                                                                                                                                                                                                                    The invention relates to a composition comprising a biologically active TR3-specific binding agent (I) that binds to TR3 and inhibits the proliferation of cells expressing TR3. (I) identified by the methods are useful for treating a subject suspected of having a disease associated with a proliferation of cells expressing TR3 especially leukemias or lymphomas or a T-cell mediated disease especially autoimmune diseases such as myasthenia gravis, systemic lupus erythamatosus, rheumatoid arthritis, diabetes, multiple sclerosis, sarcoidosis, myocarditis, thyroiditis and tumours. (I) is also useful for treating a subject suspected of having graft-versus-host disease, rejection of a transplanted organ such as heart, liver, lung, kidney, pancreas, bowel, skin or an appendage, or inflammatory diseases, allergies and contact dermatitis. The present sequence represents the nucleotide sequence of
                                                        1008
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Composition for treatment of T-cell mediated disease e.g. arthritis, cancer comprises a biologically active TR3-specific binding agent especially a monoclonal antibody -
                                                                                                                                                                                                                                                                                           Sequence 1250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 72; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tittle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          P-PSDB; AAB84941.
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                        21
GluAlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGlu
                                                 ValMetAspAlaValProAlaArgArgTrpLysGluPheValArgThrLeuGlyLeuArg
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deletion which alters the reading frame"
/transl_except= "(pos:558.559, aa:Cys)"
/note= "this codon has an apparent 1 nucleotide
deletion which alters the reading frame"
/transl_except= "(pos:638.639, aa:Leu)"
/note= "this codon has an apparent 1 nucleotide
deletion which alters the reading frame"
/transl_except= "(pos:718.719, aa:Met)"
/note= "this codon has an apparent 1 nucleotide
deletion which alters the reading frame"
/transl_except= "(pos:718.719, aa:Met)"
/note= "this codon has an apparent 1 nucleotide
deletion which alters the reading frame"
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73..1247
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Alignment Scores: Pred. No.:

Sequence 1254

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT89427
                                                                                                                                                                 This cDNA clone codes for human death domain containing receptor DR3 (see AAW31517), a novel member of the tumour necrosis factor receptor family. It was isolated from a HUVEC cDNA library. Related death domain containing receptor DR3-V1 cDNA (see AAT89426)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1128
                                                  produce recombinant polypeptides in transformed host cells. These polypeptides can be used to treat diseases and disorders associated with the inhibition of apoptosis. Antagonistscan be used to treat
                                                                                                                            was isolated from a human testis tumour cDNA library. The g
have also been identified in cDNA libraries of foetal liver,
foetal brain, tonsil and leukocyte. Nucleic acids encoding
                                                                                                                                                                                                                                             Claim
                                                                                                                                                                                                                                                                                  Death domain containing receptor polypeptide(s) DR3 and DR3-V1 for activation of apoptosis and NF-kappaB, antagonists can be
                                                                                                                                                                                                                                                                                                                                                                         Dillon
                                                                                                                                                                                                                                                                       to treat inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-0CT-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1188 GAGCGCATGGGGCTGGACGGCTGCGTGGAAGACTTGCGCAGCCGCCTGCAGCGCGGCCCG
                                                                                           foetal brain, tonsil and leukocyte. Nucleic acids encoding full-length or mature DR3, or the extracellular, transmembrane, intraclelular or especially the death domain of DR3, can be used to
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                                                                                                                                                                                                                                                                                                                                                                                                                 (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        61 GluArgMetGlyLeuAspGlyCysValGluAspLeuArgSerArgLeuGlnArgGlyPro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              41 MetLeuLysArgTrpArgGlnGlnGlnProAlaGlyLeuGlyAlaValTyrAlaAlaLeu
                                                                                                                                                                                                                                                                                                                                1997-470812/43.
DB; AAW31517.
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                                                                                                                                                                                                                                             6;
                                                                                                                                                                                                                                                                                                                                                                      ₽J,
                                      inhibition of apoptosis. Antagonistscan be used to transl disorders associated with increased apoptosis and
                                                                                                                                                                                                                                          Page 75-77; 108pp; English.
                         inflammatory diseases and disorders
                                                                                                                                                                                                                                                                                                                                                                        Dixit VM,
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 NF-kappaB; ds.
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              The invention provides nucleotide sequences encoding death domain containing receptor polypeptides DR3-V1 and DR3. The DR3-V1 cDNA clone is contained in ATCC deposition No. 97456 and the DR3 cDNA clone is contained in ATCC deposition No. 9757. Recombinant vectors comprising the nucleic acid sequences and optionally the leader sequences are
                                                                                                                                                                                                                                    WPI;
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17-OCT-1996;
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96US-0013285.
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                                                                                                                                                                                                              Human; death domain containing receptor; DR3; cancer; autoimmune disorder; inflammation; cardiovascular disorder;
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28-MAY-1999;
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                                 2000-687263/67
)B; AAB36265.
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) NI J.
/) GENTZ R L.
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Treating graft-versus-host disease, cancer, immunodeficiency autoimmune disease comprising administering an antibody to De Containing Receptor proteins and a second therapeutic agent .

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Best Local Similarity:
Query Match:
DB:
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(GETH ) GENENTECH INC
                      23-SEP-1996;
01-APR-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      arrhythmias, recurrence, neurodegenerative diseases included and congenital heart defects, neurodegenerative disease such as multiple Alzheimer's and Parkinson's diseases, autoimmune disease such as multiple sclerosis, arthritis, diabetes, Graves' disease, asthma and psoriasis, and to promote and openesis and wound healing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention provides the protein and coding sequences for two death domain containing receptors, designated DR3 and DR3-V1. These receptors are involved in apoptosis, and the sequences given can be used in the treatment of cancers, infections, cardiovascular disorders such as arrhythmias, ischaemia, aneurysms, arterial occlusive diseases, embolisms and congenital heart defects, neurodegenerative diseases including
                                                                                                                                                                                                                                                          Homo
                                                                                                                                                                                                                                                                                 Apo-3; apoptosis;
                                                                                                                                                                                                                                                                                                         Human apoptosis protein Apo-3 cDNA clone FH20.57
                                                                                                                                                                                                                                                                                                                                     14-APR-1998
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                                                           HOMO
                                                                                      Rheumatoid arthritis;
                                                                                                                   Human genomic DNA encoding
                                                                                                                                                   15-AUG-2001
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                                                         sapiens.
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                                                                                                                     a rheumatoid arthritis associated
                                                                                                                                                                                                                 ВP
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                                                                                        human;
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Location/Qualifiers 89..1342

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RESULT 7
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DT 22-A
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KW Huma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1100
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transmembrane
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)B; AAB97370.
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                                                                                                                                 rheumatoid arthritis associated
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Matches:
Conservative:
Mismatches:
Indels:
                                                                                     antirheumatic;
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                                                                                                                                 DR3 gene related
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Query Match:
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                                                                                                                                                                                                                                                                                                                      Percent Similarity:
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27-MAR-2001;
30-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to the human DR3 gene, which is associated with rheumatoid arthritis. Certain mutations in the gene can be linked to the disease. The sequences can be used to evaluate disease onset and its possibility and to provide therapy and remedies. The present sequence is a coding sequence described in the exemplification of the invention.
                                                                                                                                                 1220
                                                                                                                                                                                    Genomes, particularly DR3 genomic DNA, participating arthritis via mutation, useful in evaluating disease possibility and providing therapy and remedies -
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                                                AAC91477;
                                                                   AAC91477
                                                                                                         1280 GAGCGCATGGGGCTGGACGGCTGCGTGGAAGACTTGCGCAGCCGCCTGCAGCGCGCCCG
                                                                                                                                                                                                                                                                                                                                                                       Sequence 1634 BP;
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                                                                                                                   GluArgMetGlyLeuAspGlyCysValGluAspLeuArgSerArgLeuGlnArgGlyPro
                                                                                                                                                         MetLeuLysArgTrpArgGlnGlnGlnProAlaGlyLeuGlyAlaValTyrAlaAlaLeu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2002-417132/44.
DB; AAO17879.
                                                                                                                                               ATGCTCAAGCGCTGGCGCCAGCAGCAGCCCGCGGGCCTTCGGAGCCGTTTACGCGGCCCTG
                                                                                                                                                                                                                                                                                                            Similarity:
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SHIOZAWA S.
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2001JP-0090546.
2001JP-0099990.
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Indels:
Gaps:
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Matches:
Conservative:
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                                                                                                                                                                                                                                                                                                                                                                       0 other;
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21-MAR-2001

(first entry)

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disorders such as systemic lupus erythematosus, rheumatoid arthritis, costeoarthritis, juvenile chronic arthritis, spondyloarthropathies, systemic sclerosis, idiopathic inflammatory myopathies, Sjogren's systemic vasculitis, sarcoidosis, autoimmune haemolytic anaemia, autoimmune thrombocytopaenia, thyroiditis, diabetes mellitus, immune-mediated renal disease, demyelinating diseases of the central immune-mediated renal disease, demyelinating diseases of the central and peripheral nervous systems (such as multiple sclerosis, idiopathic demyelinating polyneuropathy or Guillain-Barre syndrome, and chronic inflammatory demyelinating polyneuropathy), hepatobiliary diseases (such as infectious, autoimmune chronic active hepatitis, primary biliary cirrhosis, granulomatous hepatitis and sclerosing cholangitis), inflammatory bowel disease, gluten-sensitive enteropathy and Whipple's disease, autoimmune or immune-mediated skin diseases (such as bullous skin diseases, erythema multiforme, contact dermatitis, psoriasis),
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01-SEP-1999
15-SEP-1999
15-SEP-1999
29-OCT-1999
30-NOV-1999
01-DEC-1999
00-DEC-1999
00-DEC-1999
00-DEC-1999
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antiallergic;
hepatobiliary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence is one of thirty three nucleic acids encoding PRO polypeptides. The PRO polypeptides, anti-PRO antibodies, agonists and antagonists are useful for treating and diagnosing immune related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Thirty three nucleic acids encoding PRO polypeptides which are useful in the diagnosis and treatment of immune related disorders, e.g. systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Wood
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antirheumatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
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20-JUL-1999;
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DB; AAB50918.
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99WO-US211547

99WO-US21547

99US-0162566

99WO-US28634

99WO-US28634

99US-0170262

99WO-US30911

2000WO-US00376

2000WO-US00376

2000WO-US04341

2000WO-US04341

2000WO-US04341

2000WO-US048414

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99US-0144758
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Kabakoff
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ive; hepatotropic; virucide;
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RESULT 9
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AC ABK4
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AC CDNA
XX Huma
AK Huma
KW Huma
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KW gene
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Best Local Similarity:
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108-MAR-1999
111-MAR-1999
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12-JUN-1999
22-JUN-1999
22-JUN-1999
22-JUL-1999
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26-JUL-1999
27-JUL-1999
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11-SEP-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   allergic diseases such as asthma, allergic rhinitis, atopic diseases food hypersensitivity and urticaria), immunological diseases lung (such as eosinophilic pneumonias, idiopathic pulmonary f and hypersensitivity pneumonitis), transplantation associated including graft rejection and graft-versus-host diseases.
                                                                                                                                                                                               11-FEB-2000;
                                                                                                                                                                                                                                                                                                                           Human; PRO; benign tumour;
                                                                                                                                                                                                                                                                                                                                                 cDNA encoding human PRO779
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                                                                                                                                                                                                                                                                                                    leukaemia; neuronal di
inflammatory disorder;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard;
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                                                                                                                                                                                                                                                                                        cytostatic; neuroprotective; gene; ss
99US-146222P.
99US-149395P.
99US-151689P.
99WO-US20111.
99WO-US21190.
99WO-US28301.
99WO-US28301.
                                                                                       99WO-USO5028
99US-13972P.
99US-133459P.
99WO-US12252.
99US-140650P.
99US-140653P.
99US-144758P.
99US-145698P.
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413.00
100.008
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disorder; stromal disorder; blastocoelic
er; immune disorder; angiogenic disorder;
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Matches:
Conservative:
Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                             lymphoid malignancy;
er; blastocoelic disorder;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polypeptides and the polynucleotide sequences encoding them. The PRO polypeptides, agonists, antagonists or anti-PRO antibodies are useful for treating benign or malignant tumours (e.g. renal, kidney, bladder, breast, etc), leukaemias and lymphoid malignancies, other disorders such as neuronal, glial, astrocytal, hypothalamic, glandular, macrophagal, stromal and blastocoelic disorders, inflammatory, immune and angiogenic disorders. The polynucleotide sequences are also useful in gene therapy. ABK40254-ABK40288 encode for the human PRO polypeptides of the invention.
                                                                                                                                                                                                                1294
                                                                                                                                                                                                                                                               1234
                                                                                                                                                                                                                                                                                                             1174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Thirty five nucleic acids encoding PRO polypeptides, useful for treating benign or malignant tumours, leukaemias and lymphoid malignancies, inflammatory, angiogenic and immunologic disorders
                                                                     Death domain
                                                                                                                   02-MAR-1998
                                                                                                                                                                                                                                                                                                                                                         1114 GTGATGGACGCGGTCCCAGCGCGCGCGCTGGAAGGAGTTCGTGCGCACGCTGGGGCTGCGC
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                                                                                                                                           AAT89426;
                                                                                                                                                                  AAT89426 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to the
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                                                         inflammation;
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                                                                                                                                                                                                                                                                                                                         21 GluAlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGlu
                                                                                                                                                                                                                                                                                                                                                                    1 ValMetAspAlaValProAlaArgArgTrpLysGluPheValArgThrLeuGlyLeuArg
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DB; AAU86139.
                                  sapiens
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                                                                                                                                                                                                                                                            ATGCTCAAGCGCTGGCGCCAGCAGCAGCCCGCGGGCCTCGGAGCCGTTTACGCGGCCCTG
                                                                                          domain
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                                                       containing; NF-kappaB;
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Pan J, P
Wood WI;
                                                                                         containing receptor DR3-V1 cDNA
                                                                                                                  (first entry)
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                                                         receptor; DR3-V1; human; apoptosis;
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i RM, Roy MA,
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Stone DM;
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RESULT 11

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GluArgMetGlyLeuAspGlyCysValGluAspLeuArgSerArgLeuGlnArgGlyPro 80

GAGCGCATGGGGCTGGACGGCTGCGTAGAAGACTTGCGCAGCCGCCTGCAGCGCCGGCCCG

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Best Local Similarity: Query Match:
DB:

2.91e-44 413.00 100.00% 100.00% 100.00%

Mismatches: Indels: Gaps:

Length: Matches: Conservative:

1783 80 0 0 0

Alignment Scores: Pred. No.: Percent Similarity:

Score:

US-09-993-234-6\_COPY\_338\_417 (1-80) x AAT89426 (1-1783)

1 ValMetAspAlaValProAlaArgArgTrpLysGluPheValArgThrLeuGlyLeuArg

GTGATGGACGCGGTCCCAGCGCGCGCGCTGGAAGGAGTTCGTGCGCACGCTGGGGCTGCGC

GluAlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGlu

1361

40 1301 20

1421 60

MetLeuLysArgTrpArgGlnGlnGlnProAlaGlyLeuGlyAlaValTyrAlaAlaLeu GAGGCAGAGATCGAAGCCGTGGAGGTGGAGATCGGCCGCTTCCGAGACCAGCAGTACGAG

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domain containing receptor DR3-VI (see AAW31516), a novel member of the tumour necrosis factor receptor family. It was isolated from a cDNA library derived from cells of a human testis tumour. Related death domain containing receptor DR3 cDNA (see AAR89427) was isolated from a HUVEC cDNA library. The genes have also been identified in cDNA libraries of foetal liver, foetal brain, tonsil and leukocyte. Nucleic acids encoding full-length or mature DR3-VI can be used to produce recombinant polypeptides in transformed host cells. These polypeptides can be used to treat diseases and disorders associated with the inhibition of apoptosis. Antagonists, such as antibodies raised against DR3-VI, can be used to treat diseases and disorders associated with increased apoptosis and for treating inflammatory
                                                                                                                                                                                                                                                                               Death domain containing receptor for activation of apoptosis and we to treat inflammatory diseases
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 Sequence 1783 BP;
                                                                                                                                                                                                                                                       Claim
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                                                                                                                                                                                                                       cDNA clone, deposited as ATCC 97456,
                                                                                                                                                                                                                                                    2; Page 71-73; 108pp; English.
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                              and
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                              disorders.
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304..1481
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 330 A; 562
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                                                                                                                                                                                     codes for human death w31516), a novel member of y. It was isolated from a
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                                                                                                                                                                                                                                containing receptor polypeptides DR3-V1 and DR3. The DR3-V1 cDNA clone is contained in ATCC deposition No. 97456 and the DR3 CDNA clone is contained in ATCC deposition No. 97757. Recombinant vectors comprising the nucleic acid sequences and optionally the leader sequences are used for the recombinant production of the proteins. The present sequence represents a cDNA encoding the death domain containing receptor polypeptide (DR3-V1).
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12-MAR-1996;
17-OCT-1996;
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                                                                                                                                                                                                                       Sequence 1783
                                                                                                                                                                                                                                                                                                                                      Claim
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                                                                                  1 ValMetAspAlaValProAlaArgArgTrpLysGluPheValArgThrLeuGlyLeuArg
MetLeuLysArgTrpArgGlnGlnGlnProAlaGlyLeuGlyAlaValTyrAlaAlaLeu
                              GAGGCAGAGATCGAAGCCGTGGAGGTGGAGATCGGCCGCTTCCGAGACCAGCAGTACGAG
                                         GluAlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGlu
                                                                      GTGATGGACGCGGTCCCAGCGCGCGCGCTGGAAGGAGTTCGTGCGCACGCTGGGGCTGCGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     domain;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   receptor; DR3-V1; DR3; recombinant; ds
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96US-0013285.
96US-0028711.
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198..300
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                                                                                                                                                                                                                                                                                              The present invention provides the protein and coding sequences for two death domain containing receptors, designated DR3 and DR3-V1. These receptors are involved in apoptosis, and the sequences given can be used in the treatment of cancers, infections, cardiovascular disorders such as arrhythmias, ischaemia, aneurysms, arterial occlusive diseases, embolisms and congenital heart defects, neurodegenerative diseases including
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Treating graft-versus-host disease, cancer, immunodeficiency or an autoimmune disease comprising administering an antibody to Death Decontaining Receptor proteins and a second therapeutic agent -
                                                                                                                                                                                           Sequence 1783
                                                                                                                                                                                                                                                     and congenital heart defects, neurodegenerative diseases including Alzheimer's and Parkinson's diseases, autoimmune disease such as musclerosis, arthritis, diabetes, Graves' disease, asthma and psorian and to promote and transfer of the content and promote and the promo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Fig 1;
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28-MAY-1999;
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) DILLON P J.
) DIXIT V M.
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                                                                                                                                                                                                                                    pound
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                                                                                                                                                                                         564 G;
                        Conservative: Mismatches: Indels:
                                                                                             Length: Matches:
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                                                                                                                                                                                                                                    healing.
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US-09-993-234-6\_COPY\_338\_417 (1-80)

ValMetAspAlaValProAlaArgArgTrpLysGluPheValArgThrLeuGlyLeuArg

x AAC68776 (1-1783)

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Percent Similarity:
Best Local Similarity:
                                 Alignment Scores: Pred. No.: Score:
                                                                                                                               This sequence encodes the human apoptosis inducing receptor (AIR) of the invention. AIR is a Type I transmembrane protein, soluble forms of which can be used to regulate cell death in a therapeutic setting. Soluble AIR can also be used in vitro to block apoptosis or AIR-expressing cells, or to screen agonists or antagonists of AIR activity. The cytoplasmic domain of AIR can be used to develop assays for inhibitors of AIR-induced cell death, which is useful to regulate cell death in a therapeutic setting as well as in vitro. Agonists of AIR activity can be used to kill tumour cells that express AIR, or T cells expressing AIR in autoimmune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1422
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Apoptosis inducing receptor; AIR protein; human; cell death regulator; Type I transmembrane protein; tumour cell death; autoimmune disease;
                                                                                                                                                                                                                                                                                                            Claim
                                                                                                                                                                                                                                                                                                                                          DNA encoding apoptosis inducing receptor - which is Type transmembrane protein, useful for regulating cell death
                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1998-240077/21.
P-PSDB; AAW57045.
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Perkins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                     Sequence 1847 BP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2; Page 28-30; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   96US-0044456.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         inducing receptor coding sequence
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                                                                                                   316 A; 605 C;
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                 exon
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                                                                                                                     exon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; DR3; rheumatoid arthritis; antirheumatic; antiarthritic;
gene therapy; gene; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1307
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                                                                                                                                                                                                                                                                                                                            exon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAL47185;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               intron
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human DR3 gene associated with rheumatoid arthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21 GluAlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGlu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 ValMetAspAlaValProAlaArgArgTrpLysGluPheValArgThrLeuGlyLeuArg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GluArgMetGlyLeuAspGlyCysValGluAspLeuArgSerArgLeuGlnArgGlyPro 80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MetLeuLysArgTrpArgGlnGlnGlnProAlaGlyLeuGlyAlaValTyrAlaAlaLeu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    therapy; gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GAGCGCATGGGGCTGGACGGCTGCGTGGAAGACTTGCGCAGCCGCCTGCAGCGCGCCCG
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   3069..3124
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Query Match:
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27-MAR-2001;
30-MAR-2001;
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                                                                                                                                                 Sequence 4825
                                                                                                                                                                                                                                    Genomes, particularly DR3 genomic DNA, participating in rheumatoid arthritis via mutation, useful in evaluating disease onset and its possibility and providing therapy and remedies -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           exon
                                                                                                                                                                 with rheumatoid arthritis. the disease. The sequences possibility and to provide the gene of the invention.
                                                                                                                                                                                                     The
                                                                                                                                                                                                                    Claim 1; Page 64-66; 84pp; Japanese.
                                                                                                                                                                                                                                                                         WPI; 2002-417132/44.
                                                                                                                                                                                                                                                                                         Shiozawa S,
                                                                                                                                                                                                                                                                                                         (NEWI-)
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                                             L
                                                                                                                                                                        present invention relates to the human DR3 gene, which is associated h rheumatoid arthritis. Certain mutations in the gene can be linked to disease. The sequences can be used to evaluate disease onset and its sibility and to provide therapy and remedies. The present sequence is
                          GluAlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGlu
                                                                                                                                 Scores:
                                                                                                                                                                                                                                                                                                         NEW IND RES SHIOZAWA S.
                                                                                                                                                                                                                                                                                                                                  ; 2000JP-0324296.
; 2001JP-0090546.
; 2001JP-0099990.
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3530..3577
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3578..4021
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3334..3529
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      14-AUG-2000
12-AUG-2000
22-AUG-2000
22-AUG-2000
22-AUG-2000
23-AUG-2000
21-SEP-2000
01-SEP-2000
01-SEP-2000
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07-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human;
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17-MAR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Genomic sequence #589 encoding novel human connective tissue polypeptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          connective tissue related disorder;
atic; gene; ds.
                                         2000US-0226868.
2000US-0227182.
2000US-0227009.
2000US-0228924.
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                                                                                                                                                       Nucleic acid encoding novel connective tissue associated polypeptides, used in diagnosing, preventing, treating or ameliorating a disorder such as cancer or rheumatoid arthritis -
                                                                                                                                                                                                WPI;
                                                                                                                                      Disclosure;
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The present invention relates to the isolation of novel human connective tissue related polypeptides (AAU86435-AAU86923) and the polynucleotide (cDNA and genomic) sequences encoding them. The sequences of the invention are useful in the diagnosis, treatment, prevention and/or prognosis of diseases associated with connective tissue(s), including cancer. The polynucleotide sequences of the invention are also useful in gene therapy. ABK42102-ABK43116 represent genomic sequences encoding the novel human connective tissue related polypeptides.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published\_pct\_sequences. 2003 3215

Sequence 10797 BP; ۶, **;** 3378 ပ္ 2201 T; 0 other;

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Page 12

Search completed: April 6, 2003, 16:52:44 Job time: 81.489 secs

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SUMMARIES

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Submitted (19-MAR-1997) Molecular Immunology
Molecular Medicine, John Radcliffe Hospital,
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LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing
PTOC. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
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Accession Number U94501, probable skipping of putative
exons 3, 4, 5, 6 and 7"
236 c 248 g 109 t
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LFCCRGCPADEAGMEALTPPPATHLSPLDSAHTLLAPPDSSEKICTVQLVGNSWTPGY
PETQEALCPQVTWSWDQLPSRALGPAAAPTLSPESPAGSPAMMLQPGPQLYDVMDAVP
ARRWKEFVRTLGLREAEIEAVEVEIGRFRDQQYEMLKRWRQQQPAGLGAVYAALERMG
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/protein_id="AAC51313.1"
/db_xref="GI:2071963"
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/note="similar to Fas and TNF-R1;
/not
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/db_xref="taxon:9606"
/cell_line="HeLa"
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HSU94507
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                                                                                       Query Match:
DB:
                                                  US-09-993-234-6_COPY_338_417 (1-80) x HSU94507 (1-838)
                                                                                                                         Best Local Similarity:
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1 ValMetAspAlaValProAlaArgArgTrpLysGluPheValArgThrLeuGlyLeuArg
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LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing
Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human lymphocyte a spliced, complete U94507
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (19-MAR-1997) Molecular Immunology Group, Institute of Molecular Medicine, John Radcliffe Hospital, Headington, Oxford
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Direct Submission
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                                                                                                                                                                                                                                                                exon 3"
328. .329
                                                                                                                                                                                                                                        /note-"deletion compared to LARD-1a, deposited in Accession Number U94501, probably due to skipping putative exons 5, 6 and 7, leading to premature transcriptional termination"

281 c 285 g 138 t
                                                                                                                                                                                                                                                                                                                                                                                                                               /product="lymphocyte associated receptor of death 6"
/protein_id="AAC51312.1"
/db_xref="Gi:2071961"
/translation="MEQRPRGCAAVAAALLLVLLGARAQGGTRSPRCDCAGDFHKKIG
LFCCRGCPAASQVALENCSAVADTRCGCKPGWFVECQVSQCVSSSPFYCQPCLDCGAL
                                                                                                                                                                                                                                                                                                                                                             /note="deletion compared to LARD 1a, deposited in GenBank Accession Number U94501, probable skipping of putative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         transmembrane domain; possibly soluble; LARD-6"
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/db_xref="taxon:9606"
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596 GTGATGGACGCGGTCCCAGCGCGCGCGCTGGAAGGAGTTCGTGCGCACGCTGGGGCTGCGC

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Direct Submission
Submitted (19-MAR-1997) Molecular Immunology
Submitted (19-MAR-1997) Molecular Immunology
Molecular Medicine, John Radcliffe Hospital,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Screaton,G.R., Xu,X.N., Olsen,A.L., Cowper,A.E., Tan,R., McMichael,A.J. and Bell,J.I.
LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
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                                                                                   407. .408

Anotes deletion compared to LARD-la, deposited in Accession Number U94501, probably due to skipping putative exons 6 and 7, leading to premature transcriptional termination to 317 c 319 g 162 t
                                                                                                                                                                                      Accession Number
                                                                                                                                                                                                                              /translation="MEQRPRGCAAVAAALLLVLLGARAQGGTRSPRCDCAGDFHKKIG
LFCCRGCPAASQVALENCSAVADTRCGCKPGWFVECQVSQCVSSSPFYCQPCLDCGAL
HRHTRLLCSRRDTDCGTCLPGFYEHGDGCVSCPT"
                                                                                                                                                                                                                                                                     /product="lymphocyte associated receptor of death
/protein_id="AAC51311.1"
/db_xref="GI:2071959"
                                                                                                                                                                                                                                                                                                                           soluble; truncated before the
                                                                                                                                                                                                                                                                                                                                       /function="mediates apoptosis"
/note="LARD-5; similar to Fas
                                                                                                                                                                                                                                                                                                                                                                                           /organism="Homo sapiens"
/db_xref="taxon:9606"
/cell_line="HeLa"
                                                                                                                                                                                      /note="deletion compared
Accession Number U94501,
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US-09-993-234-6_COPY_338_417 (1-80) x HSU94506 (1-952)
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LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing
PTOC. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
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Submitted (19-MAR-1997) Molecular Immunology
Molecular Medicine, John Radcliffe Hospital,
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                                                                          /note="deletion compared to LARD la, deposited in GenBank Accession Number 094501, probable skipping of putative
                                                                                                                                                                                    /product="lymphocyte associated receptor of death 4"
/protein_id="AAC51310.1"
/protein_id="AAC51310.1"
/db_xref="G1:2071957"
/tanslation="MEQREPRECANVANALLLYLLGARAQGGTRSPRCDCAGDFHKKIG
LFCCRGCPAGHYLKAPCTEPCGNSTCLVCPQDTFLAWENHHNSECARCQACDEQASQV
ALENCSAVADTRCGGKPGWFYECQVSQCVSSSPFYCQPCLDCGALHRHTRLLCSRRDT
                                                                                                                                                                                                                                                                                                                                                                            /function="mediates apoptosis"
/note="LARD-4; similar to Fas
soluble; truncated before the
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/db_xref="taxon:9606"
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Submitted (19-MAR-1997) Molecular Immunology Group, Institute
Molecular Medicine, John Radcliffe Hospital, Headington, Oxfor
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LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
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         /translation="MEQRPRGCAAVAAALLLVLLGARAQGGTRSPRCDCAGDFHKKIG LFCCRGCPAASQVALENCSAVADTRGGCKPGWFVECQVSQCVSSSPFYCQPCLDCGAL HRHTRLLCSRRDTDCGTCLPGFYEHGDGCVSGPTSCGSCPERCAAVCGWRQMFWQQV LLAGLVVFLLLGATLTYTYHKCWFHKFLVTADEAGMEALTPPPATHLSPLDSAHTLLA PPDSSEKICTVQLVGNSWTPGYPTGEALCPQVTWSWDQLPSRALGPAAAPTLSPESPAGSPAMMLQPGPQLXDVMDAVPARKWKEFVRTLGLREAEIEAVEVEVEIGRFRDQQYEML
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KRWRQQQPAGLGAVYAALERMGLDGCVEDLRSRLQRGP'
                                                                                                /codon_start=1
/product="lymphocyte associated
/protein_id="AAC51314.1"
/db_xref="GI:2071965"
                                                                                                                                                              TNF-R1"
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/note="LARD-8; NGFR family member;
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/db_xref="taxon:9606"
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U94510
U94510.1 GI:2071966
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                                                                                                                                                                                                                                                                                                                                                                        Screaton,G.R., Xu,X.N., Olsen,A.L., Cowper,A.E., Tan,R., McMichael,A.J. and Bell,J.I.
LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing
PTOC. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
                                                                                                                                                                                                                                                       Submitted (19-MAR-1997) Molecular Immunology Group, Institute of Molecular Medicine, John Radcliffe Hospital, Headington, Oxford OX3
                                                                                                                                                                                                                                                                                                             2 (bases 1 to 1143)
Screaton, G.R.
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                                                                                                                                                                                                                                                                                              Direct Submission
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/note="deletion compared the Accession Number U94501, person 3"
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/codon_start=1
/product="lymboyte associated receptor of death
/protein_id="AAC51315.1"
/db_xref="G1:2071967"
                                                                  TNF-R1"
                                                                                                                                      /ce11
                                                                                                                                                                    /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                        Location/Qualifiers
                                                                                /function="mediates
/note="LARD-9; NGFR
                                                                                                                                                    cell_line="HeLa"
                                                                                                                  -11_Type="lymphocyte"
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probable skipping of putative
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                                                                                                                                                                                                                                   Proc. Nat
97272273
                                                                                                                                                                                                                                           Screaton,G.R., Xu,X.N., Olsen,A.L., Cowper,A.E., Tan,R., McMichael,A.J. and Bell,J.I.
LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing
Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
                                                                                                                 you,
                                                                                                                             Submitted (19-MAR-1997) Molecular Immunology Group, Institute of Molecular Medicine, John Radcliffe Hospital, Headington, Oxford, OX3
                                                                                                                                                                                                                                                                                                                                                                                       Homo
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                                                                                                                                                                  Direct Submission
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                                                                                                                                                                                   Screaton, G.R.
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PTLSPESPAGSPAMMLQPGPQLYDVMDAVPARRWKEFVRTLGLREAEIEAVEVEIGRF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /translation="MEQRPRGCAAVAAALLLVLLGARAQGGTRSPRCDCAGDFHKKIG
LFCCRGCPAGHYLKAPCTEPCGNSTCLVCPQDTFLAWENHHNSECARCQACDEQASQV
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DCGTCLPGFYEHGDGCVSCPTSTLGSCPERCAAVCGWRQNEAGMEALTPPPPATHLSPL
                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                    Eutheria; Primates;
/cell_type="lymphocyte"
l. .657
                               /cell_line="HeLa"
                                             /organism="Homo sapiens"
/db_xref="taxon:9606"
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AX150176
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 Similarity:
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1 (bases 1 to 1250)
Tittle, T.V. and Wegmann, K.W.
Tr3-specific binding agents and methods
patent: WO 0135995-A 1 25-MAY-2001;
Tittle, Thomas V. (US); Wegmann, Keith
Location/Qualifiers
                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                     Sequence 1
AX150176
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LFCCRGCPAGHYLKAFCTEPGCNSTCLVGPQDTFLAWERHHUSSECARCOACDEQASQV
ALENCSAVADTRCGGCREMYYECQVSSQVSSSPFYCOPCLDCGALHRHFRLLCSSRDF
DCGTCLPGFYEHGDGCVSCPTVLGPGAPGWPCGPPPAMGHPDLHIPPLLASQAPGYCR
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soluble; truncated before
                                                                                                                /organism="Homo sapiens"
/db_xref="taxon:9606"
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/db_xref="GI:2071955"
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Sequence 3
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     Human death receptor U72763
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Yu.G.-L., Ni,J., Gentz,R.L. and Dillon,
Death domain containing receptors
Patent: US 6153402-A 3 28-NOV-2000;
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                                              HSU78029
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/product="death receptor 3"
/protein_id="acath receptor 3"
/db_xref="GI:1669512"
/db_xref="GI:
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Submitted (12-NOV-1996) Biochemistry,
University St., Seattle, WA 98101, USA
                                      Human lymphocyte associated receptor
                                                         HSU94501
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Mammalia; Eutheria;
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ALENCSAVADTRCGCKPGMETVEQVSQCVSSSPTYQDCLDCGALHRHIFALLGSRDT
DCGTCLPGFYEHGDGCVSCPTSTLGSCPERCAAVCGWRQMFWVQVLLAGLVVPLLLGA
TLTYTYTXHCWPHKFLVVTADEAGWEALTPPPATHLSFLSAFTLLAFDDSSEKICTVQL
VGNSWTPGYPETQEALCPQVTWSWDDLPSRALGPAAPTLSPESPAGSFMANDAPPA
UGDVDADAPPARKETYFTLGLREAFIEAVEVEIGRERDQQYEMLKRWRQQQPAGLGA
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/protein_id="AAB40918.1"
/db_xref="GI:1778764"
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/db_xref="taxon:9606"
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                                                                   1192 GAGCGCATGGGGCTGGACGGCTGCGTGGAAGACTTGCGCAGCCGCCTGCAGCGCCGGCCCG
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                                                                                                                                                      41 MetLeuLysArgTrpArgGlnGlnGlnProAlaGlyLeuGlyAlaValTyrAlaAlaLeu 60
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Submitted (19-MAR-1997) Molecular Immunology Group, Institute
Molecular Medicine, John Radcliffe Hospital, Headington, Oxfor
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VYAALERMGLDGCVEDLRSRLQRGP"
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ALENCSAVADTRCGCKPGWFVECQVSQCVSSSFFYCQPCLDCGLHRHTRLCSSRDT
DCGTCLPGFYEHGDGCVSCPTSTLGSCPERCAAVCGWRQMFWYQVLAGLVVPLLGS
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TNF-R1; contains a death
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/translation="MEQRPRGCAAVAAALLLVLLGARAQGGTRSPRCDCAGDFHKKIG
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/protein_id="AAC51306.1"
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/cell_line="HeLa"
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LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing
Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
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/protein_id="AAC51307.1"
/db_xref="GI:2071951"
/db_xref="GI:2071951"
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LFCCRGCPAGHYLKAPCTEPCGNSTCLVCPQDTFLAWENHHNSECARCQACDEQASQV
ALENCSAVADTRCGCKPGWFVECQVSQCVSSSPFYCQPCLDCGALHRHTRLLCSRRDT
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/note="LARD-lb; membrane prote
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Accession Number U94501, probably rep;
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/db_xref="taxon:9606"
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LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing
Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                      223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                lymphocyte associated receptor
                                                                                                                                                                                                                                                                /note="insertion compared to LARD la, depos
Accession Number U94501, probable retained
to premature transcriptional termination"
441 c 451 g 240 t
                                                                                                                                                                                                                                                                                                                                                    /product="lymphocyte associated receptor of death 2"
/protein_id="AAC51308.1"
/protein_id="AAC51308.1"
/db_xref="Gi:2071953"
/translation="MEQRPRGCAAVAALLLVILGARAQGGTRSPRCDCAGDFHKKIG
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GEEGGNHPTPTSCFQCSGSRCSWLALMSPSCLGPP"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="LARD-2; similar to Fas and TNF-R1; possibly soluble; truncated before the transmembrane domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               complete cds
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /codon_start=1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                function="mediates_apoptosis"
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GluAlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGlu

40

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1173 GAGGCAGAGATCGAAGCCGTGGAGGTGGAGATCGGCCGCTTCCGAGACCAGCAGTACGAG 1232

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Direct Submission
Submitted (03-JAN-1997) Molecular Biotechnology, University of
Washington, 1705 NE Pacific Street, HSB-K360, Seattle, WA 9819
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human death domain receptor 3 U83597
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Chaudhary, P.M. and Hood, L.E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                    280
                                                                                                                                                                                                                                                                                                                                                               /translation="GCAAVAAALLLVLLGARAQGTRSPRCDCAGDEHKKIGLECCRG
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LGFYEHGDGCVSCPTSTLGSCPERCAAVCGWRQMFWVQVLLAGLVVPLLLGATLTYTY
RHCWPHKPLVTADEAGHEALTPPPATHLSPLDSAHTLLAPPDSSEKICTVQLVGNSWT
PGYPETQEALCPQVTWSWDQLPSRALGPAAAPTLSPESPAGSPAMMLQPGPQLYDVMD
                                                                                                                                                                                                                                                                                                                 RMGLDGCVEDLRSRLQRGP"
496 c 470 g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note="A TNER1-related death-domain containing receptor; DR3, WSL-LR, Apo-3, TRAMP; transmembrane form; similar to human EST clone 298913, GenBank Accession Number N71143"
                                                                                                                                                                                                                                                                                                                                                AVPARRWKEFVRTLGLREAEIEAVEVEIGRFRDQQYEMLKRWRQQQPAGLGAVYAALE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /product="death domain receptor 3"
/protein_id="AAB41432.1"
/db_xref="GI:1800293"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'gene="DDR3"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    function="mediates apoptosis when cross-linked"
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Search completed: April 6, 2003, 21:08:02 Job time: 933.72 secs \E\ |

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Result
No.
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-MODEL-frame+_p2n.model -DEV=xlh
-Q-/cgn2_1/USPTQ_spool/US09993234/runat_27032003_115457_15378/app_query.fasta_1.2346
-Q-/cgn2_1/USPTQ_spool/US09993234/runat_27032003_115457_15378/app_query.fasta_1.2346
-DB-Issued_Patents_NA -QFMT-fastap -SUFFIX=rn1 -MIMMATCH=0.1 -LOOPCL=0
-LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi
-LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15
-MODE=LOCAL -OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=200000000
-USER=-US0999324_eCGN_1_1_116_erunat_27032003_115457_15378 -NCPU=6 -ICPU=3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
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1: /cgn2_6/ptodata//
2: /cgn2_6/ptodata//
3: /cgn2_6/ptodata//
4: /cgn2_6/ptodata//
5: /cgn2_6/ptodata//
6: /cgn2_6/ptodata//
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length: 2000000000
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Ygapop 10.0 , Y
Fgapop 6.0 , F
Delop 6.0 , I
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                                                                                                                                                                                                                                                                                                                                                                                                                                  /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
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US-08-928-069-11

US-08-828-683A-9

US-08-815-469-1

US-08-762-338-10

US-09-513-007-1

US-09-397-787-236

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US-09-106-038A-1
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1799 3881 2268 2040 2040 2040 2040 2015 2152 2152 2152 2152 2152 2152 215	2175 2175 2175 2175 2413 1062 1236 1323 1799
<b>44124444444413443342348</b> 04	114C44W4
US-09-079-029-2 US-09-33-593A-1 US-08-444-005-14 US-08-53-669A-5 US-09-183-861-5 US-09-183-861-5 US-09-183-861-1 US-09-185-918-1 US-09-48-868-1 US-09-48-868-1 US-09-16-443-1 US-09-16-443-1 US-09-16-443-1 US-09-18-944-005-16 US-09-18-944-005-16 US-09-18-944-005-16 US-09-18-944-005-16 US-09-18-94-1-1 US-09-18-94-1-1 US-09-18-94-1-1 US-08-868-1-1 US-08-868-1-1 US-08-954-668-1-1	-08-837 -08-126 -08-054 -08-651 -09-333 -09-134 -08-883 -09-329
Sequence 2, Appli sequence 14, Appli sequence 5, Appli sequence 5, Appli sequence 5, Appli sequence 2, Appli sequence 2, Appli sequence 1, Appli sequence 1, Appli sequence 16, Appli sequence 17, Appli sequence 2, Appli sequence 2, Appli sequence 2, Appli sequence 3, Appli sequence 7, Appli sequence 7, Appli sequence 7, Appli sequence 18, Appli	

## ALIGNMENTS

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US-08-815-469-3
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Patent No. 6153402
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                                                                                                                                                           COMPUTER: IBM PC COMPACTIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/815,469
FILING DATE: HEREWITH
CLASSIFICATION: 435
                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: No. 1
FILING DATE: 06-FEB-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 66
                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Yu, Guo-Liang
APPLICANT: Ni, Jian
APPLICANT: Dixit, Vishva
APPLICANT: Gentz, Reiner L.
FILING DATE: 17-OCT-PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Dillon, Patrick J.
TITLE OF INVENTION: Death Domain Containing Receptors
NUMBER OF SEQUENCES: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 1100 New CITY: Washington
                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
ZIP: 20005-3934
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1100 New York Ave.,
                                        17-OCT-1996
  us 60/013,285
                                                                                                                        6153402 Yet
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FILING DATE: 12-MAR-1996 ATTORNEY/AGENT INFORMATION:

Steffe,

Eric K.

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Query Match:
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                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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                                                                         MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech) CURRENT APPLICATION DATA:
CURRENT AFFLICATION NUMBER: US/08/920, APPLICATION 11-Sep-1997
FILING DATE: 11-Sep-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 60/026943
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1012
                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 inc
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
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CITY: S
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STRANDEDNESS: double
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4FR: 1488.0310003/EKS/KRM
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Matches:
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Patent NO. 6469144
GENERAL INFORMATION:
APPLICANT: Ashkenazi,
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INFORMATION FOR SEQ ID NO:
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                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/625328
FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb:
MEDIUM TYPE: 15.5 inch, 1.44 Mb:
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                        SOFTWARE: WinPatin (CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Apo-2 LI AND Apo-3 POLYPEPTIDES NUMBER OF SEQUENCES: 28
                                    ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Genente
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                                                                                                                                               APPLICATION NUMBER: US/08/828,683A FILING DATE: 31-Mar-1997 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                   CITY: South San F: STATE: California
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                                                                                                                                                                                                                                                                                                               ZIP: 94080
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                   Marschang,
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Diane L.
BER: 35,600
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US-08-815-469-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1,
APPLICATION NUMBER: US/08/815,469
FILING DATE: HEREWITH
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: No. 6153402 Yet
FILING DATE: 06-FEB-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/028,711
FILING DATE: 17-OCT-1996
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                                                                                                                                                 COMPUTER READABLE FORM:
MEDIIM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
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APPLICANT: Ni, Jian
APPLICANT: Dixit, V
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APPLICANT: Gentz, Reiner L.
APPLICANT: Dillon, Patrick J.
TITLE OF INVENTION: Death Domain Containing Receptors
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                                                                                                                                                                                                                                                  20005-3934
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  , Application 6153402
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1100 New York Ave.,
                                                                                                                                                                                                                                                                     USA
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                                                                     6153402 Yet Assigned
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; LOCATION:
US-08-815-469-1
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                                                                                                                                                                                                                                                                                              Sequence 10, Application US/08762308 Patent No. 5925548
                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 12-MAR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Steffe, ETIC K.
            SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                  1362
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                      TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                        1422 GAGCGCATGGGGCTGGACGGCTGCGTGGAAGACTTGCGCAGCCGCCTGCAGCGCGGCCCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1302 GAGGCAGAGATCGAAGCCGTGGAGGTGGAGATCGGCCGCTTCCGAGACCCAGCAGTACGAG
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                                                                                                                                                                                                                                                 APPLICANT: Beutler, APPLICANT: Bazzoni,
                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                 61 GluargMetGlyLeuaspGlyCysValGluaspLeuargSerargLeuGlnargGlyPro 80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21 GluAlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGlu
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TYPE: nucleic acid
STRANDEDNESS: double
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                                                                                                                                                               STREET:
                                                                                                                  COUNTRY:
APPLICATION NUMBER:
                                                                                                                                                                             ADDRESSEE:
                                                                                                                                               Houston
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us/08/762,308
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Matches:
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Indels:
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CLASSIFICATION: 530 PRIOR APPLICATION DATA:

FILING DATE:

APPLICATION NUMBER: US OF FILING DATE: 05-APR-1994

US 08/224,593

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                                                                                     SEQ ID NO
                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                            Sequence 1, Applic Patent No. 6406907
                                                                                                                                      APPLICANT: Mwangi, Simon
TITLE OF INVENTION: BOVINE TUMOR NECROSIS FACTOR RECEPTOR-1
TITLE OF INVENTION: AND METHODS OF USE
FILE REFERENCE: 08411-018001
CURRENT APPLICATION NUMBER: US/09/513,007
CURRENT FILING DATE: 2000-02-25
PRIOR APPLICATION NUMBER: 60/122,156
PRIOR APPLICATION NUMBER: 60/122,156
PRIOR FILING DATE: 1999-02-26
                                                                                                                                                                                                                                                                                                 APPLICANT: Taylor, J. Michael APPLICANT: Kehrli, Jr., Marcus
                                                                                                        SOFTWARE:
                                                                                                                         NUMBER OF SEQ ID NOS:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                     APPLICANT:
                               TYPE: DNA
ORGANISM: Bos taurus
NAME/KEY: CDS
                  FEATURE:
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REFERENCE/DOCKET NUMBER: UTS
TELECOMUNICATION INFORMATION:
TELEPHNOE: 418-3000
TELEPHNI: 474-777-77
                                                                     ENGTH: 2440
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STRANDEDNESS: sing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GluAlaGluIleGluAlaValGluValGluIleGlyArg---PheArgAspGlnGlnTyr 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GTGGTGGATGGCGTGCCTCCAGCGCGCTGGAAGGAGTTCATGCGTTTCATGGGGGCTGAGC 1238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TyrAlaAlaLeuGluArgMetGlyLeuAspGlyCysValGluAspLeuArgSerArgLeu 76
                                                                                                      FastSEQ for Windows Version
                                                                                                                                                                                                                                                                                                                                                                           Application US/09513007
                                                                                                                                                                                                                                                                                    Lee, Eun-Kyung
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; SEQ ID NO 236
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; TYPE: DNA
; ORGANISM: Homo sapien
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Best Local Similarity:
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US-09-513-007-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Mitcham, Jennifer L.
APPLICANT: King, Gordon E.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR OVARIAN
TITLE OF INVENTION: CANCER THERAPY AND DIAGNOSIS
FILE REFERENCE: 210121.466C2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/397,787 CURRENT FILING DATE: 1999-09-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1605 GGCCGCGTGCTCAGGGACATGGACCTGCTGGGTTGCCTGGAAAACATAGAGGAGGCGCTG 1664
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                                                                     113
                       40 GluMetLeuLysArgTrpArgGlnGlnGlnPro------AlaGlyLeuGlyAlaVal 56
                                                                                                                                  53 GTGGTGGAGAACGTGCCCCCGTTGCGCTGGAAGGAATTCGTGCGGCGCCCTAGGGCTGAGC 112
                                                                                             21 GluAlaGluIleGluAlaValGluValGluIleGlyArg---PheArgAspGlnGlnTyr 39
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                                                              AGCATGCTGGCGACCTGGAGGCGGCGCGCACGCGCGCGAGGCCACGCTGGAGCTGCTG 232
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Best Local Similarity:
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (415) 327-32 INFORMATION FOR SEQ ID NO:
                                                                                                                                                           1238 GTGGTGGAGAACGTGCCCCCGTTGCGCTGGAAGGAATTCGTGCGGCGCCTAGGGCTGAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 2062 base pairs
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TITLE OF INVENTION: Modified human TNFalpha (Tumor
TITLE OF INVENTION: Necrosis Factor alpha) Receptor
                                            FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
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                                                                                                                               21 GluAlaGluIleGluAlaValGluValGluIleGlyArg---PheArgAspGlnGlnTyr 39
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CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 51
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                                                                                                                                                                                      1 ValMetAspAlaValProAlaArgArgTrpLysGluPheValArgThrLeuGlyLeuArg
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GluMetLeuLysArgTrpArgGlnGlnGlnPro-----AlaGlyLeuGlyAlaVal 56
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15) 327-3231
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Best Local Similarity:
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; LOCATION:
US-08-465-982-24
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Patent No. 5863786
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INFORMATION FOR SEQ ID NO:
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                                   1358 AGCATGCTGGCGACCTGGAGGCGGGGGCGCACGCGGGGGGGCGAGGCCACGCTGGAGCTGCTG 1417
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ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, version #1.25 CURRENT APPLICATION DATA:
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                                                                                                       MOLECULE TYPE: FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: RObbins, Roberta L.
REGISTRATION NUMBER: 33,208
REFERENCE/DOCKET NUMBER: 5150-0030
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
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57 TyrAlaAlaLeuGluArgMetGlyLeuAspGlyCysValGluAspLeuArgSerArgLeu 76
                                                                                                                                          21 GluAlaGluIleGluAlaValGluValGluIleGlyArg---PheArgAspGlnGlnTyr 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                     GluMetLeuLysArgTrpArgGlnGlnGlnPro-----AlaGlyLeuGlyAlaVal 56
                                                                                                                                                                              GTGGTGGAGAACGTGCCCCCGTTGCGCTGGAAGGAATTCGTGCGGCGCCCTAGGGCTGAGC 1297
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US-09-505-250-3
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Query Match:
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US-09-106-038A-1
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 Patent
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INFORMATION FOR SEQ ID NO:
                                                                              1519
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CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Laurel Spear Bernstein
NAME: James NUMBER: 37,280
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MEDIUM TYPE: 3.5 inch disk,
COMPUTER: IBM PC compatible
                                                                                                                                                                                                        1339 GTGGTGGAGAACGTGCCCCCGTTGCGCTGGAAGGAATTCGTGCGGCGCCCTAGGGCTGAGC 1398
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TELECOMMUNICATION INFORMATION:
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CORRESPONDENCE ADDRESS:
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ce 3, Application US/09505250A
No. 6329148
                                                                                                                                                              GluMetLeuLysArgTrpArgGlnGlnGlnPro-----AlaGlyLeuGlyAlaVal 56
                                                                           GGACGCGTGCTCCGCGACATGGACCTGCTGGGCTGCCTGGAGGACATCGAGGAGGCGCTT 1578
                                                                                                       TyralaalaLeuGluargMetGlyLeuAspGlyCysValGluaspLeuArgSerArgLeu 76
                                                                                                                                         AGCATGCTGGCGACCTGGAGGCGGCGCGCGCGCGCGCGAGGCCACGCTGGAGCTGCTG
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VENTION: ANTISENSE MODULATION
VENTION: EXPRESSION
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TITLE OF INVENTION: Synergistic Anti-Cancer Therapy with
TITLE OF INVENTION: Triptolides and Death Domain Ligands
FILE REFERENCE: SUN-109PRV2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: SUN-109FRV2
CURRENT APPLICATION NUMBER: US/09/505,250A
CURRENT FILING DATE: 2000-02-15
NUMBER OF SEQ ID NOS: 4
                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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LOCATION: (25
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                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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                                         SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: BROWDY AND NEIMARK
                                                                                                                                                                                                                                                                                                               APPLICANT:
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                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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            APPLICATION NUMBER: US/0 FILING DATE: 12-OCT-1994
                                                                                                                                                             COUNTRY:
                                                                                                                                                                                STATE:
                                                                                                                                                                                             CITY: Washington
                                                                                                                                                                                                                STREET:
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CLASSIFICATION:
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                                                                                                                                                                                                             419 Seventh Street, N.W.,
                                                                                                                                                                                                                                                                                                              BATKIN, Michael
                                                                                                                                                             USA
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                             US/08/321,668
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ATTORNEY/AGENT INFORMATION:

PRIOR APPLICATION DATA:
APPLICATION NUMBER: IL 1
FILING DATE: 12-OCT-1993

IL 107268

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APPLICANT: WALLACH
APPLICANT: BRAKEBU
APPLICANT: BATKIN,
APPLICANT: BATKIN,
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 2175 base pair
COMPUTER READABLE FORM:
MEDIOM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
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                                                                                                                                                                                                       TITLE OF INVENTION:
                                                                                                                                                                                                                     APPLICANT: BATKIN, Michael TITLE OF INVENTION: MOLECU
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REGISTRATION NUMBER: 25,618
REFERENCE/DOCKET NUMBER: WAI
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                                                                                              COUNTRY:
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VARFOLOMEEV, Eugene
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Matches:
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US-08-126-016-1
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Best Local Similarity:
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                                                                                                                                                                                                   Patent No. 581126
                                                                                                                                                                                    GENERAL INFORMATION:
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                                                                                                                                        APPLICANT:
APPLICANT:
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REFERENCE/DOCKET NUMBER: WAI
TELECOMMUNICATION INFORMATION:
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LENGTH: 2175 base pairs
TYPE: nucleic acid
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                                       NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                   APPLICANT: ADERKA, DAN TITLE OF INVENTION: EXP
                                                                                                                                                                                                                                                                          1:339
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STREET:
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                           ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                GluMetLeuLysArgTrpArgGlnGlnGlnPro-----AlaGlyLeuGlyAlaVal 56
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          E: Browdy and Neimark
419 Seventh Street, N.W.,
                                                                                                                                        NOPHAR, YARON
KEMPER, OLIVEI
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                                                                                                            KEMPER, OLIVER
ENGELMANN, HARTMUT
BRAKEBUSCH, CORD
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Matches:
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CURRENT APPLICATION DATA:

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25

COMPUTER READABLE FORM:

20004 Ŕ

USA

MEDIUM TYPE:

Floppy disk

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US-08-126-016-1
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Best Local Similarity:
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                                                                        Sequence 1, Application US/08054970 Patent No. 6395267
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                             GENERAL INFORMATION:
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LOCATION:
FEATURE:
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                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
TELEFAX: 202-737-3528
                                                                                                                                         1519 GGACGCGTGCTCCGCGACATGGACCTGCTGGGCTGCCTGGAGGACATCGAGGAGGCGCTT
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REFERENCE/DOCKET NUMBER:
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FILING DATE: 24-SEP-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                            GluMetLeuLysArgTrpArgGlnGlnGlnPro-----AlaGlyLeuGlyAlaVal 56
                                                                                                                                                                                                                                              TyralaalaLeuGluArgMetGlyLeuAspGlyCysValGluAspLeuArgSerArgLeu 76
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Search completed: April Job time: 19.5049 secs

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US-08-054-970-1
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Best Local Similarity:
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COMPUTER READABLE FORM:
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APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                      1339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 2175 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION: TELEPHONE: 202-628-5197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
                                     1519 GGACGCGTGCTCCGCGACATGGACCTGCTGGGCTGCCTGGAGGACATCGAGGAGGCGCTT 1578
                                                                                           TELLEFAX: ZUL
TELLEFAX: Z48633
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Townsend, Kevin G. REGISTRATION NUMBER: 34,033 REFERENCE/DOCKET NUMBER: WA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 0:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
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                                                                                                                 40 GluMetLeuLysArgTrpArgGlnGlnGlnPro-----AlaGlyLeuGlyAlaVal 56
                                                                                                                                                                        21 GluAlaGluIleGluAlaValGluValGluIleGlyArg---PheArgAspGlnGlnTyr 39
                                                                                                                                                                                                                   GTGGTGGAGAACGTGCCCCCGTTGCGCTGGAAGGAATTCGTGCGGCGCCTAGGGCTGAGC 1398
                                                               TyralaalaLeuGluargMetGlyLeuAspGlyCysValGluAspLeuArgSerArgLeu 76
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Matches:
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Indels:
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Sequence 9, Appli Sequence 1, Appli Sequence 1577, Ap Sequence 1389, Ap Sequence 1389, Ap

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Result
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-Q-/cgn2_l/USPTQ_spool/US09993234/runat_27032003_115459_15466/app_query.fasta_1.2346
-Q-/cgn2_l/USPTQ_spool/US09993234/runat_27032003_115459_15466/app_query.fasta_1.2346
-DB-Published_Applications_NA -QFWT-fastap -SUFFIX=rnpb -MINMATCH=0.1
-LOOPEL-0 -LOOPEXT=0 -UNITS-bits -START=1 -END--1 -MATRIX-blosum62
-TRANS-human40.cdi -LIST=45 -DCALIGN=200 -THR_SCORE-pct -THR_MAX=100
-THR_MIN=0 -ALIGN=15 -MODE=LOCAL -OUTPWT-pto -NORW-ext -HEAPSIZE=500 -MINLEN=0
-MAXLEN=200000000 -USER-US0993234_eCGN_pto -NORW-ext -HEAPSIZE=500 -MINLEN=0
-NCPU-6 -LCPU=3 -NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG
-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPOEXT=0.5 -FGAPOP=6
-FGAPOEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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Listing first 45 s
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    /cgn2_6/ptodata/1/pubpna/US07_NEM_PUB.seq:*
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US-09-333-966-3
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                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                        APPLICANT: Dixit, Vishva
APPLICANT: Gentz, Reiner L.
APPLICANT: Dillon, Patrick J.
TITLE OF INVENTION: Death Domain
NUMBER OF SEQUENCES: 17
                                                                                                                COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                APPLICANT: Yu, Guo-Liang
APPLICANT: Ni, Jian
APPLICANT: Dixit, Vishva
                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                    COUNTRY: USA
ZIP: 20005-3934
                                                                                                                                                                  CITY: Washington STATE: DC
                                                                                                                                                                                            ADDRESSEE: Sterne, Kes
STREET: 1100 New York
                               APPLICATION NUMBER:
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               FILING DATE:
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US-09-874-138-1
US-09-874-138-1
US-09-992-964-10
US-09-9828-739-1
US-09-887-879-10
US-09-887-879-10
US-09-811-088-5
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US-09-764-847-1577
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US-09-917-800A-1601
US-09-99-234-14
US-09-999-422-14
US-09-99-76-89-236-1
US-09-99-76-89-236-1
US-09-99-76-89-236-1
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US-09-89-422-16
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Sequence 14, Appl
Sequence 24, Appl
Sequence 11, Appl
Sequence 11, Appl
Sequence 10, Appl
Sequence 100, App
Sequence 1, Appl
Sequence 1, Appl
Sequence 1, Appl
Sequence 1, Appl
Sequence 16, Appl
Sequence 16, Appl
Sequence 16, Appl
Sequence 16, Appl
Sequence 17, Appl
Sequence 17, Appl
Sequence 11, App

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US-10-081-280-9
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                                                                                                                                                                             Sequence 9, Application US/10081280 Patent No. US20020165157A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 202-371-2540
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 14
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 12-MAR-1996 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 17-OCT-PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 06-FEB-PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                          41
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LOCATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: No. US20020009773A1 Yet Assigned FILING DATE: 06-FEB-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ENGTH:
ZIP: 94080
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 i
                                                                                                    APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-2 LI AND
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
                                                                                                                                                                                                                                                              GluAlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGlu 40
                                                                                                                                                                                                                                                                                                                                         MetLeuLysArgTrpArgGlnGlnGlnProAlaGlyLeuGlyAlaValTyrAlaAlaLeu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nucleic acid
                                                         STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STATE: California
                                             COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Steffe, Eric K.
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US-10-112-793-9
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                                                                                                                                                                                                                     Sequence 9, Application US/10112793
Publication No. US20020192729A1
GENERAL INFORMATION:
APPLICANT: AShkenazi, Avi J.
TITLE OF INVENTION: Apo-2 LI
NUMBER OF SEQUENCES: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: 9:
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                                                                                                                                                                                                                                                                                                                                                                                                                               61 GluargMetGlyLeuAspGlyCysValGluaspLeuArgSerArgLeuGlnargGlyPro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21 GluAlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGlu 40
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb:
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 1634 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION: TELEPHONE: 415/225-5416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech) CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATGCTCAAGCGCTGGCGCCAGCAGCAGCCCGCGGGCCTTCGGAGCCGTTTTACGCGGCCCCTG
                                                                                                                                                                                                                                                                                                                                                                                                           GAGCGCATGGGGCTGGACGGCTGCGTGGAAGACTTGCGCAGCCGCCTGCAGCGCGGCCCG
                                                                                                                    CITY: South San Francisco
STATE: California
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                                                                                  ZIP:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007R1
                                                                                                   COUNTRY:
                                                                                                                                                                STREET: 1 DNA Way
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: 910/371-7168
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                                                                                                                                                                                   ADDRESSEE:
                                                                                    94080
                                                                                                     USA
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LI AND Apo-3 POLYPEPTIDES
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Indels:
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                                   floppy
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PRIOR APPLICATION DATA:

CLASSIFICATION: <Unknown>

APPLICATION NUMBER: US/10/112,793 FILING DATE: 28-Mar-2002

CURRENT APPLICATION DATA:

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RESULT 4
US-10-112-193-11
; Sequence 11, Application US/10112193
; Publication No. US20030004313A1
; GENERAL INFORMATION:
; GENERAL INFORMATION:
AShkenazi, Avi J.
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Query Match:
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SEQUENCE CHARACTERISTICS:
LENGTH: 1634 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                     61 GluargMetGlyLeuAspGlyCysValGluAspLeuArgSerArgLeuGlnArgGlyPro 80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21 GluAlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGlu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 ValMetAspAlaValProAlaArgArgTrpLysGluPheValArgThrLeuGlyLeuArg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 08/625328
FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 650/952-9881
COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
                                                                                                                                                                                                                           APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-3 POLYPEPTIDE
NUMBER OF SEQUENCES: 15
                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MetLeuLysArgTrpArgGlnGlnGlnProAlaGlyLeuGlyAlaValTyrAlaAlaLeu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GAGGCAGAGATCGAAGCCGTGGAGGTGGAGATCGGCCGCTTCCGAGACCAGCAGTACGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATGCTCAAGCGCTGGCGCCAGCAGCAGCCCGCGGGCCTCGGAGCCGTTTACGCGGCCCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Marschang, Diane L. REGISTRATION NUMBER: 35,600
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                                                                                                                                  CITY: South San Francisco
STATE: California
                                                                                                              COUNTRY:
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                                                                                               ZIP: 94080
                                                                                                                                                                                          ADDRESSEE: Genentech,
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Indels:
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Best Local Similarity:
Query Match:
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US-09-993-234-9
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                                                                                                                                                                                                                                                                                Sequence 9, Application US/09993234
Patent No. US20020146768A1
GENERAL INFORMATION:
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TITLE OF INVENTION: Apo-2 LI AND Apo-3 POLYPEPTIDES

NUMBER OF SEQUENCES: 11

CORRESPONDENCE ADDRESS:

ADDRESSEE: Genentech, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: MAISCHANG, DIANE L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1052R1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 60/026943
FILING DATE: 23-Sep-1996
ATTORNEY/AGENT INFORMATION:
                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 1
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              CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                    GluArgMetGlyLeuAspGlyCysValGluAspLeuArgSerArgLeuGlnArgGlyPro 80
                                                                                                                                                                                                                                                                                                                                                                                       GAGCGCATGGGGCTGGACGCCTGCGTGGAAGACTTGCGCAGCCCCTGCAGCGCCCCG
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                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech)
                                                                                                                                              STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 1634 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 08/928,069 FILING DATE: 11-Sep-1997
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                                                                                                                                COUNTRY: USA
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US/09/993,234
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Matches:
Conservative:
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Indels:
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FILING DATE: 19-NO. US20020146768A1-2001 CLASSIFICATION: <Unknown>

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Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                                                                                                                         Sequence 1, Application US/09333966 Patent No. US20020009773A1
                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 9: SEQUENCE CHARACTERISTICS:
           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                                                                     APPLICANT: Yu, Guo-Liang
APPLICANT: Ni, Jian
APPLICANT: Dixit, Vishva
APPLICANT: Gentz, Reiner L.
APPLICANT: Dillon, Patrick J.
TITLE OF INVENTION: Death Domain Containing Receptors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1160 GAGGCAGAGCCGAGAGCCGTGGAGGTGGAGATCGGCCGCTTCCGAGACCAGCAGTACGAG
                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                       1280 GAGCGCATGGGGCTGGACGGCTGCGAAGACTTGCGCAGCCGCCTGCAGCCGCCGGCCCG
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                                                                                                                                                                                                           NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                        61 GluArgMetGlyLeuAspGlyCysValGluAspLeuArgSerArgLeuGlnArgGlyPro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21 GluAlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGlu
                                                                                                            COUNTRY:
                                                                                                                                                             STREET:
                                                                                                                                                                             ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-5416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: Linear SEQUENCE DESCRIPTION: SEQ
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATGCTCAAGCGCTGGCGCCAGCAGCAGCCGCGGGCCTCGGAGCCGTTTACGCGGCCCTG
                                                                                               20005-3934
APPLICATION DATA:
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TELEX: 910/371-7168
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1100 New York Ave.,
                                                                                                              USA
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                Version #1.30
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; APPLICANT: Rosen et al.; TITLE OF INVENTION: Nucleic Acids, Proteins, an; FILE REFERENCE: PC009C1; CURRENT APPLICATION NUMBER: US/10/092,154; CURRENT FILING DATE: 2002-03-07; NUMBER OF SEQ ID NOS: 2003; Prior Application removed - See File Wrapper or; SOFTWARE: Patentin Ver. 2.0; SEQ ID NO 1577
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Best Local Similarity:
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Publication No. US20030054375A1
GENERAL INFORMATION:
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FILING DATE: 12-MAR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Steffe, Eric K.
REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 1488.031
                                                                                                                                                                                                                                                                                                1422 GAGCGCATGGGGCTGGACGGCTGCGTGGAAGACTTGCGCAGCCGCCTGCAGCGCGGCCCG 1481
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LOCATION:
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STRANDEDNESS: double
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**LENGTH: 10797** 

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RESULT 8
US-09-764-847-1577
; Sequence 1577, Application US/09764847
; Patent No. US20020132767A1
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DB:
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DB:
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; ORGANISM: Homo
US-10-092-154-1577
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Best Local Similarity:
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; ORGANISM: HOMO
US-09-764-847-1577
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APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PC009
                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Ver. SEQ ID NO 1577
                                                                                                                                                                                                                                                                                                                                                                                                                                     Prior application data removed - consult PALM NUMBER OF SEQ ID NOS: 2003
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CURRENT FILING DATE: 2001-01-17
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                                            MetLeuLysArgTrpArgGlnGlnGlnProAlaGlyLeuGlyAlaValTyrAlaAlaLeu
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                             ATGCTCAAGCGCTGGCGCCAGCAGCAGCCGGGCCTCGGAGCCGTTTACGCGGCCCTG
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Patent NU. COLOR
GENERAL INFORMATION:
APPLICANT: Mendrick, Donna
APPLICANT: Porter, Mark
APPLICANT: Porter, Mark
APPLICANT: Johnson, Kory
APPLICANT: Castle, Arthur
APPLICANT: Gene Logic, Inc.
TITLE OF INVENTION: Molecular Toxicology Modeling
FILE REFERENCE: 44921-5038-US
CURRENT APPLICATION NUMBER: US/09/917,800A
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US-09-917-800A-1601
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Query Match:
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; ORGANISM: Homo
US-09-954-531-1389
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PRIOR FILING DATE: 2000-09-18
PRIOR PPLICATION NUMBER: US/60/234,009
PRIOR FILING DATE: 2000-09-20
PRIOR APPLICATION NUMBER: US/60/234,034
PRIOR FILING DATE: 2000-09-20
PRIOR APPLICATION NUMBER: US/60/234,509
PRIOR FILING DATE: 2000-09-22
PRIOR FILING DATE: 2000-09-22
PRIOR APPLICATION NUMBER: US/60/234,567
PRIOR FILING DATE: 2000-09-22
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LENGTH: 1743
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Patent No. US20020119462A1
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CURRENT FILING DATE: 2002-05-02
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APPLICATION NUMBER: US 60/303,459

APPLICATION NUMBER: US 60/298,884

APPLICATION NUMBER: US 60/297,457

2001-06-13 2001-06-19

APPLICATION NUMBER: US 60/292,336
FILING DATE: 2001-05-22
APPLICATION NUMBER: US 60/295,798
FILING DATE: 2001-06-06

FILING DATE:

FILING DATE:

APPLICATION NUMBER: US 60/290,645

2001-05-15 2001-05-11

APPLICATION NUMBER: US 60/290,029

APPLICATION NUMBER: US 60/222,880

NUMBER: US 60/222,040: 2000-07-31

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                                                                                                                                                                           Sequence 14, Application US/09898234 Patent No. US20020155112A1
                                                                                            APPLICANT:
APPLICANT:
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LENGTH: 2130
                                             TITLE OF INVENTION: THE RECEPTORS, TITLE OF INVENTION: Them FILE REFERENCE: 98,385-I
                CURRENT APPLICATION NUMBER: US/09/898,234 CURRENT FILING DATE: 2001-07-03
                                                                                                                        APPLICANT: Hauptmann, Rudolph APPLICANT: Himmler, Adolph
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APPLICATION
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SEQ ID NOS: 1740
                                                                                          Maurer-Fogy, Ingrid
Stratowa, Christian
NUMBER: 09/525,998
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Best Local Similarity:
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                                                                                                                                                                                                    APPLICANT: Stratowa, Christian TITLE OF INVENTION: TNF Receptors, TITLE OF INVENTION: Them
                                                                                                       CURRENT FILING DATE: 2001-07-03
PRIOR APPLICATION NUMBER: 09/792,356
PRIOR FILING DATE: 2000-02-23
PRIOR FILING DATE: 2000-02-23
                                                                                                                                         FILE REFERENCE: 98-385-J
CURRENT APPLICATION NUMBER: US/09/899,429A
CURRENT FILING DATE: 2001-07-03
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                                                                     PRIOR APPLICATION NUMBER: 08/477,639 PRIOR FILING DATE: 1955-06-07
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PRIOR FILING DATE: 1993-11-17
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                                 APPLICATION NUMBER: 08/383,676 FILING DATE: 1995-02-01
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               APPLICATION NUMBER: 08/153,287
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5. US20020169118A1
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APPLICATION NUMBER: 07/821,750 FILING DATE: 1992-01-02

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                                            PRIOR APPLICATION NUMBER: 08/383,676
PRIOR FILLING DATE: 1995-02-01
PRIOR APPLICATION NUMBER: 08/153,287
PRIOR FILING DATE: 1993-11-17
PRIOR APPLICATION NUMBER: 07/821,750
PRIOR FILING DATE: 1992-01-02
PRIOR APPLICATION NUMBER: 07/511,430
PRIOR FILING DATE: 1990-04-20
SEQ
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                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/792,356
CURRENT FILING DATE: 2001-08-17
PRIOR APPLICATION NUMBER: 08/477,639
PRIOR FILING DATE: 1995-06-07
                                                                                                                                                                                                                                                             TITLE OF INVENTION: THE RECEPTORS, TITLE OF INVENTION: Them FILE REFERENCE: 98,385-G
                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Hauptmann, Rudolph APPLICANT: Himmler, Adolph
                             NUMBER OF SEQ ID NOS: 87
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Stratowa, Christian
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PRIOR FILING DATE: 2000-03-15
PRIOR APPLICATION NUMBER: 08/383,676
PRIOR FILING DATE: 1995-02-01
PRIOR APPLICATION NUMBER: 08/153,287
PRIOR FILING DATE: 1993-11-17
PRIOR APPLICATION NUMBER: 07/821,750
PRIOR FILING DATE: 1992-01-02
PRIOR FILING DATE: 1992-01-02
                                                                                                                                        SOFTWARE: P
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                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/899,422
CURRENT FILING DATE: 2001-08-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: THE RECEPTORS, TITLE OF INVENTION: Them
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                                                                                                                                                                                                  PRIOR FILING DATE:
                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 07/511,430
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  FEATURE:
NAME/KEY: CDS
LOCATION: (245)..(1630)
OTHER INFORMATION: Desc
                                                                             ORGANISM: Artificial Sequence
                                                                                                    TYPE: DNA
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OTHER INFORMATION:
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Stratowa, Christian
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US-09-899-422-14

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GENERAL INFORMATION:

APPLICANT: Taylor, J. Michael

APPLICANT: Kehrli, Jr., Marcus

APPLICANT: Lee, Eun-Kyung

APPLICANT: Lee, Eun-Kyung

APPLICANT: Lee, Eun-Kyung

APPLICANT: Mwangi, Simon

FILE OF INVENTION: BOVINE TUMOR NECROSIS FACTOR RECEPTOR-1

FILE REFERENCE: 08411-018001

CURRENT APPLICATION NUMBER: US/09/970,532

CURRENT FILING DATE: 2001-10-03

PRIOR APPLICATION NUMBER: 09/513,007

PRIOR APPLICATION NUMBER: 09/513,007
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Search completed: April 7, 2003, 03:58:57 Job time: 41.4618 secs

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GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
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Database: EST:*  1: em_estba:* 2: em_esthum:* 3: em_estin:* 4: em_esttmu:* 5: em_esty:* 6: em_esty:* 7: em_estro:* 9: gb_est:* 10: gb_est:* 11: gb_htc:* 12: gb_est:* 13: gb_est:* 14: gb_est:* 15: em_estfun:* 16: em_estom:* 16: em_estom:* 17: gb_gss:* 18: em_gss_hum:* 19: em_gss_lnv:* 20: em_gss_pin:* 21: em_gss_pin:* 24: em_gss_num:* 25: em_gss_num:* 26: em_gss_nus:* 27: em_gss_nus:* 27: em_gss_pro:* 27: em_gss_pro:*	Command line parameters:  -MODEL-frame+ p2n.model -DEV-xlh -O-/cgn2_1/USPTO_spoo1/US09993334/runat_27032003_115456_15362/app_query.fasta_1.2346 -O-/cgn2_1/USPTO_spoo1/US09993334/runat_27032003_115456_15362/app_query.fasta_1.2346 -DB-EST -OFMY-fastap -SUFFIX-rst -MINMATCH-0.1 -LOOPCL-0 -LOOPEXT-0 -UNITS-bits -START-1 -END1 -MATRIX-blosum62 -TRANS-human40.cd1 -LIST-45 -DOCALIGN-200 -THR_SCORE-pct -THR_MAX-100 -THR MIN-0 -ALIGN-15 -MODE-LOCAL -OUTEWT-pto -NORM-ext -HEAPSIZE-500 -MINLEN-0 -MAXIEN-200000000 -USER-US0993234_@CGN_1_1_4749_@runat_27032003_115456_15362 -NCPU-6 -ICPU-3 -NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES-0 -WAIT -LONGLOG -DEV_TIMEOUT-30 -WARN_TIMEOUT-30 -THREADS-1 -XGAPPOP-10 -XGAPEXT-7 -YGAPDP-10 -YGAPEXT-0.5 -DELOP-6 -DELEXT-7	Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries	Minimum DB seq length: 0 Maximum DB seq length: 2000000000	Total number of hits satisfying chosen parameters: 32308132	Searched: 16154066 seqs, 8097743376 residues	Scoring table: BLOSUM62  Xgapop 10.0 , Xgapext 0.5  Ygapop 10.0 , Ygapext 0.5  Fgapop 6.0 , Fgapext 7.0  Delop 6.0 , Delext 7.0	Title: US-09-993-234-6_COPY_338_417  Perfect score: 413  Sequence: 1 VMDAVPARRWKEFVRTLGLRERMGLDGCVEDLRSRLQRGP 80	Run on: April 6, 2003, 16:19:54; Search time 500.409 Seconds (without alignments) 2589.161 Million cell updates/sec	OM protein - nucleic search, using frame_plus_p2n model	copyright (c) 1993 - 2003 compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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BM694442 BM694442 BM767536 BG035257 BM742388 BM742388 AU117362 BQ880824 BQ228387 BM555065 BM923204	BE138968 BF138948 BI525375 BF118096 AA631757 AW355430 AA08232 BT168232	AA088350 AW492480 AW492480 AI703436 BM009354 BE014705 AA887388	AI266746 BM794760 BM794760 BB212432 BF552058 W71984 W71984 BF726557 BE563566 AA524052 AW13906	BM695193 BM666370 BE670189 AW074008 AI380959 AI1811528 BF655336 BF655336 AW182875 AI380900 BM824360 BM824360	ID
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## ALIGNMENTS

AUTHORS TITLE	REFERENCE	SOURCE ORGANISM	KEYWORDS	VERSION	ACCESSION		DEFINITION	LOCUS	BM695193	RESULT 1
Bonaldo, M.F., Lennon, G. and Soares, M.B. Normalization and subtraction: two approaches to facilitate gene	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  1 (bases 1 to 510)	human. Homo sapiens	EST.	BM695193.1 GI:19008451	BM695193	UI-E-CO1-aev-i-03-0-UI 5', mRNA sequence.	1 Homo sa	BM695193 510 bp mRNA linear EST 28-FEB-2002		

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                                                                  181 GAGCGCATGGGCCTGGACGCCTGCGAAGACTTGCGCAGCCGCCTGCAGCGCGCCCG
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                                                                                        61 GluArgMetGlyLeuAspGlyCysValGluAspLeuArgSerArgLeuGlnArgGlyPro 80
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Genome Res.
97044477
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Tissue Procurement: Dr. Gregg Hageman
cDNA Library preparation: Dr. M. Bento Soares, Univeristy of Iowa
cDNA Library Arrayed by: Dr. M. Bento Soares, Univeristy of Iowa
DNA Sequencing by: Dr. M. Bento Soares, Univeristy of Iowa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        University of Iowa 1451 Eckstein Medical Research Building Iowa City, IA 52242, Tel: 319 335 8250 Fax: 319 335 9565
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Clone Distribution: Researchers may obtain clones from Research Genetics (www.resgen.com).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Soares, MB
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/lab_host="bhildh (Life Technologies) (Tl phage resistant)"
/note="Organ: eye; Vector: pT7T3-Pac (pharmacia) with a modified polylinker; Site_1: Ecor I; Site_2: Not I;
UI-E-CQl is a normalized cDNA library containing the following tissue(s): optic nerve. The library was constructed according to Bonaldo, Lennon and Soares, Genome Research, 6:791-806, 1996. First strand cDNA synthesis was primed with an oligo-df primer containing a Not I site. Double stranded cDNA was ligated to an EcoR I adaptor, digested with Not I, and cloned directionally into pT7T3-Pac vector. The oligonucleotide used to prime the synthesis of first-strand cDNA contains a library tag sequence that is located between the Not I site and the (df)18 tail. The sequence tag for this library is CCATPAAGTG. This library was created for the program, Gene Discovery in the Visual System, supported by National Eye Institute (NEI)."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /tissue_type="optic nerve"
/dev_stage="adult"
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/clone_lib="UI-E-CQ1"
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Query Match:
US-09-993-234-6_COPY_338_417 (1-80) x BM666370 (1-550)
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Genetics (www.resgen.com).
The following repetitive elements were sequence: 1-37, >POLY_A#Simple_repeat (m
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Tissue Procurement: Dr. Gregg Hager
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      451 Eckstein Medical Research Building Iowa City, IA 52242, USA
Tel: 319 335 8250
Fax: 319 335 9565
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50 bp mRNA linear EST 27-UI-E-CQ1-aev-i-03-0-UI.S1 UI-E-CQ1 Homo sapiens cDNA clone UI-E-CQ1-aev-i-03-0-UI 3', mRNA sequence.

BM666370
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                                                                                                                                                                                                                                                                                                                                                                                                                           synthesis was primed with an oligo-dT primer containing a Not I site. Double stranded CDNA was ligated to an EcoR I adaptor, digested with Not I, and cloned directionally into pT7T3-Pac vector. The oligonucleotide used to prime the synthesis of first-strand cDNA contains a library tag sequence that is located between the Not I site and the (dT)18 tail. The sequence tag for this library is CCATTAAGTG. This library was created for the program, Gen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /lab_hosi="DH10B (Life Technologies) (T1 phage resistant)"
/note="Organ: eye; Vector: pT7T3-Pac (Pharmacia) with a
modified polylinker; Site 1: EcoR I; Site 2: Not I;
UI-E-CQ1 is a normalized cDNA library containing the
following tissue(s): optic nerve. The library was
constructed according to Bonaldo, Lennon and Soares,
Genome Research, 6:791-806, 1996. First strand cDNA
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                                                                                                                                                                                                                                                                                                                    TAG_TISSUE=human optic nerve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,
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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cDNA Library Preparation: M. Bento Soares, Ph.D. cDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Clone distribution: NCI-CGAP clone distribution information
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                                                                                                                                                                                                                                                                                               /note="Organ: lung; Vector: pT7T3D-Pac (Pharmacia) with modified polylinker; Plasmid DNA from the normalized library NCI_CGAP_LU5 was prepared, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from a pool of 5,000 close the process of the process o
                                                                                                                                                                                                          Soares and M. Fatima Bonaldo.
215 c 225 g 134 t
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/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone_lib="NCI_CGAP_Lu24"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  www-bio.llnl.gov/bbrp/image/image.html
Seq primer: -40UP from Gibco
High quality sequence stop: 383.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tissue Procurement: Chris Moskaluk, M.D., Ph.D., Michael R. Emmert-Buck, M.D., Ph.D. cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: Christa Prange, The I.M.A.G.E. Consortium DNA Sequencing by: Washington University
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Site_1: Sall; Site_2: Notl;
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Technologies."
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/db_xref="taxon:9606"
/clone="IMAGE:2575504"
/clone_lib="NCI_CGAP_GUI"
/tissue_type="2 pooled high-grade transitional cell
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CDNA Library Preparation: M. Bento Soares, Ph.D.

CDNA Library Preparation: M. Bento Soares, Ph.D.

CDNA Library Preparation: M. Bento Soares, Ph.D.

CDNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 1237 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tumor Gene Index
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Ash Alizadeh, John Byrd, M.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 587)
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                                                                                                                                                                                       T 3'); double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and Cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is normalized, and was constructed by Bento
                                                                                                                                                                                                                                                                                                 Soares and M.Fatima Bonaldo."
192 c 176 g 120 t
                                                                                                                                                                                                                                                                                                                                                                                                             /tissue_type="B-cell, chronic lymphotic leukemia"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone="IMAGE:2109120"
/clone_lib="NCI_CGAP_CLL1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ocation/Qualifiers
           3.67e-41
399.00
        Length: Matches:
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DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D.,
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 623)
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Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT
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/lab_host="DH10B"
                                                                                                                                                           11538-014
                                                                                                                                                                                    Average insert size 1.75 kb.
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/clone_lib="NCI_CGAP_Ut1"
/tlssue_type="well-differentiated endometrial
                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/db_xref="taxon:9606"
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279811 MARC 3BOV Bos taurus cDNA
BF655336
BF655336.1 GI:11920468
                                                                                                                                                                                                                                                                                                                           Email: smith@email.marc.usda.gov
Single pass sequencing. Bases called and alt_trimmed with phred
v0.980904.e. Vector identified by cross_match with the -minscorv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Smith, T.P.L., Grosse, W.M., Freking, B.A., Roberts, A.J., Stone, R.T., Casas, E., Wray, J.E., White, J., Cho, J., Fahrenkrug, S.C., Bennett Gasas, E., Holton, M.P., Laegreid, W.W., Rohrer, G.A., Chitko-McKown, C.G., Pertea, G., Holt, I., Karamycheva, S., Liang, F., Quackenbush, J. and
                                                                                                                                                                                                                            Seq
                                                                                                                                                                                                                                            BACKWARD: GTTTTCCCAGTCACGACG Plate: 76 row: G column: 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Smith TPL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 libraries and construction of a gene index for cattle Genome Res. 11 (4), 626-630 (2001)
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                                Library made from pooled tissue from marrow, alveolar macrophage, ovary, fetal semitendonosus muscle, and fetal longissimus muscle."
                                                                                                                                    /organism="Bos taurus"
/db_xref="taxon:9913"
/clone_lib="MARC 3BOV"
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/lab_host="DH10B"
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                                                                                                                                                                                                                                                                                                                                                               Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL;
IMAGE Consortium (info@image.llnl.gov) for further i
Seq primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
National Cancer Institute, Cancer Genome Ana
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                                            /tissue_type="tumor"
/lab_host="DH10B (phage-resistant)"
/lab_host="DH10B (phage-resistant)"
/note="Organ: cervix; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; 1st strand cDNA was primed with a Not I - oligo(dT) primer [5]
                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                              /clone_lib="Soares_NHCeC_cervical_tumor"
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Length: Matches:

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                                                                                                                                                                                                                                                                                                                                                                                                                     M.D., Louis M. Staudt, M.D., Ph.D.

CDNA Library Preparation: M. Bento Soares, Ph.D.

CDNA Library Preparation: M. Bento Soares, Ph.D.

CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llni.gov/bbrp/image/image.html
Insert Length: 1246 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AI380900 544 bp mRNA linear EST 30-MAR-1999 tg17f01.x1 NCI_CGAP_CLL1 Homo sapiens cDNA clone IMAGE:2109049 3' similar to TR:000278 LYMPHOCYTE ASSOCIATED RECEPTOR OF DEATH
                                                                                                                                                                                                                                                                                                                                                                 Seg primer: -40UP from Gibco
High quality sequence stop: 348.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Ash Alizadeh, John Byrd, M.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert Strausberg, Ph.D.
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                                                                                                                   adaptors (Pharmacia), digested with Not I the Not I and Eco RI sites of the modified Library is normalized, and was constructed Soares and M.Fatima Bonaldo."
                                                                                                                                                                                                                                                               /db_xref="taxon:9606"
/clone="IMAGE:2109049"
/clone_lib="NCI_CGAP_CLL1"
                                                                                                                                                                                                                          /tissue_type="B-cell, chronic lymphotic leukemia"
/lab_host="DH10B"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kim, N.S., Hahn, Y., Oh, J.H., Lee, J.Y., Ahn, H.Y., Chu, M.Y., Kim, Oh, K.J., Cheong, J.E., Sohn, H.Y., Kim, J.M., Park, H.S., Kim, S. a
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Kim.N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EST.
                                                                                                                                                                                                                                                                                                                            Email: yongsung@mail.kribb.re.kr
Plate: 85 row: D column: 08
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (2002)
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                                                                                                                                                                                                                                                                                                              quality sequence stop: 439.
/note="Organ: Stomach; Vector: pT7T3-Pac; Site_1: EcoRI; Site_2: NotI; The S22SNUI6 library was contributed by the Soares laboratory and it was constructed as described by Bonaldo, M.F., Lennon, G. and Soares, M.B. (1996), Genome Research 6(9): 791-806. RNA was prepared from harvested cells of SNU-16 culture. SNU-16 cell was obtained from Korean Cell Line Bank (KCLB). SNU-16 was established from ascitic fluids of Korean patients by Park J.G. et al.
                                                                                                                                               /cell_type="Lymphoblast-like"
/cell_line="SNU-16"
                                                                                                                                                                                                /sex="
                                                                                                                                                                                                                              /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="S22SNU16n1-85-D08"
                                                                                                                                 /lab_host="DH10B"
                                                                                                                                                                                /tissue_type="Ascites"
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499 bp mRNA linear EST 27-MAR-1UI-H-COO-aqq-a-09-0-UI.sl NCI_CGAP_Sub9 Homo sapiens cDNA clone IMAGE:3104895 3', mRNA sequence.
                                                                                                                                                                                                                                                                                                                      CDNA Library preparation: Dr. M. Bento Soares, University of Iowa CDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa DNA Sequencing by: Dr. M. Bento Soares, University of Iowa Clone Distribution: Clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tumor Gene Index Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     National Cancer Institute, Cancer
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/lab_host="DH10B (Life Technologies)"
/note="Vector: pT773-Pac (Pharmacia) with a modified.
/note="Vector: pT773-Pac (Pharmacia) with a modified.
polylinker; Site_1: EcoR I; Site_2: Not I; tissues:
Cholonic mucosa with Crohns disease, Cholonic mucosa with
ulcerative colitis, Fetal thymus, Cervix, Cervical
adenosquamous carcinoma, Ligament cells, Prostate
carcinoma, Bladder carcinoma, Brain oligodenroga;
NCI_CGAP_Sub9 is a subtracted cDNA library constructed
                                                                                                                                            /db_xref="taxon:9606"
/clone="IMAGE:3104895"
/clone_lib="NCI_CGAP_Sub9"
/tissue_type="mixed"
/dev_stage="mixed"
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148 c 150 g
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                                                                                                                             9712 Medical Center Dr.,
Tel: 301 838 3528
Fax: 301 838 0208
                                                                                                                                                                                Contact: John Quackenbush
The Institute for Genomic Research
                                                                                                                                                                                                                 Assessment of gene expression patterns in a model of colon tumor metastasis using a 19,200 element cDNA microarray Unpublished (2000)
                                                                                                                                                                                                                                                                                            Hegde, P., Qi, R., Abernathy, K., I.E., Saeed, A.I., Sharov, V., 1
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/organism="Homo sapiens"
/db_xref="taxon:9606"
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Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Al266746 RNA linear qq11c06.x1 Soares_NhHMPU_S1 Homo sapiens cDNA clone y similar to TR:000278 000278 LWAPHOCYTE ASSOCIATER PROPERTY [2] TR:000280 ; mRNA sequence.
                                                                                                                                                                                                                                                                                                                       This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Insert Length: 837 Std Error: 0.00
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A1266746.1 GI:3884904
EST.
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National Cancer Institute, Cancer Genome Anat
                                                                                                                                                                                                                                                                                                                                                                          Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota;
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               /note="Organ: mixed (see below); Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site1: Not I; Site2: Eco RI; Equal amounts of plasmid DNA from three normalized libraries (melanocyte 2NbHM, pregnant uterus NbHPU, and fetal heart NbHHJ9W) were mixed, and scircles were made in vitro. Following Hap purification, this DNA was used as tracer in a subtractive hybridization
                                                                                                                                                       /tissue_type="Pooled human melanocyte, fetal heart,
pregnant uterus"
     reaction.
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/note="Vector: pBluescriptSKm"
155 c 156 g 111 t
                                                                                                                                                                                                      /clone="IMAGE:1932202"
                                                                                                                                                                                                                        /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                     'lab_host="DH10B"
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Tel: +82-42-860-4470
Fax: +82-42-860-4409
                                                                                                                                                                                                                                                                                                                                                  Kim, Y.S.
21C Frontier Korean
Unpublished (2002)
                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 647) (bases 1 to 647) (kim,N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,Oh,K.J., Cheong,J.E., Sohn,H.Y., Kim,J.M., Park,H.S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5', mRNA sequence.
BM794760
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                                                                                                                                                                                                                                                                                                                          Genome Research Center
                                                                                                                                                                                                                                                                                                                                        Contact: Kim YS
                                                                                                                                                                                                                                                                                         Korea Research Institute of Bioscience & Biotechnology 52 Eoeun-dong Yuseong-gu, Daejeon 305-333, South Korea
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                                                                                                                                                                                                                                                yongsung@mail.kribb.re.kr
 /note="Organ: Stomach; Vector: pT7T3-Pac; Site_1: EcoRI;
Site_2: NotI; The S22SNU16 library was contributed by th
Soares laboratory and it was constructed as described by
                                                               /tissue_type="Ascites"
/cell_type="Lymphoblast-like"
/cell_line="SNU-16"
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/clone="S22SNU16n1-48-G11"
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Arakawa,T., Carninci,P., Fukuda,S., Furuno,M., Hanagaki,T., nass,...
Arakawa,T., Carninci,P., Ishii,Y., Ito,M., Kawai,J., Konno,H., Kouda
, Hiramoto,K., Hori,F., Ishii,Y., Ito,M., Kawai,J., Konno,H., Kouda
,M., Koya,S., Matsuyama,T., Miyazaki,A., Nomura,K., Ohno,M.,
Okazaki,Y., Okido,T., Saito,R., Sakai,C., Sakai,K., Sano,H., Sasaki
,D., Shibata,K., Shinagawa,A., Shiraki,T., Sogabe,Y., Suzuki,H.,
,D., Shibata,K., Shinagawa,A., Takeda,Y., Tanaka,T., Toya,T.,
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BB212432 RIKEN full-length enriched, 0 day neonat musculus cDNA clone A430110122 3', mRNA sequence.
     prepare full-length cDNA libraries for rapid discovery of new genes. Genome Res. 10 (10), 1617-1630 (2000) waq1.K., Fullwake c room.
                                                    Email: genome-res@gsc.riken.go.jp,
URL:http://genome.gsc.riken.go.jp/
Carninci,P., Shibata,Y., Hayatsu,N., Sugahara,Y., Shibata,K., Itoh
,M., Konno,H., Okazaki,Y., Muramatsu,M. and Hayashizaki,Y.
Normalization and subtraction of cap-trapper-selected cDNAs to
                                                                                                                                                                                                              Laboratory for Genome Exploration Research Group, RIKEN Genomic Sciences Center(GSC), Yokohama Institute
The Institute of Physical and Chemical Research (RIKEN)
1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Jap
Tel: 81-45-503-9222
                                                                                                                                                                                                                                                                                                                                  On Jun 30, 2000 this sequence version replaced gi:8877385.
Contact: Yoshihide Hayashizaki
                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (2001
                                                                                                                                                                                                                                                                                                                                                                                                                                  Tagami,M., Tagawa,A., Takahashi,F., Takeda,Y.,
Muramatsu,M. and Hayashizaki,Y.
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41 MetLeuLysArgTrpArgGlnGlnGlnProAlaGlyLeuGlyAlaValTyrAlaAlaLeu 60

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                              21 GluAlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGlu 40
                                                                                            1 ValMetAspAlaValProAlaArgArgTrpLysGluPheValArgThrLeuGlyLeuArg
CDNA library was prepared and sequenced in Mouse Genome
Encyclopedia Project of Genome Exploration Research Group in
Genomic Sclences Center and Genome Sclence Laboratory in RIKE
Division of Experimental Animal Research in Riken contributed
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Mapping of 19032 mouse cDNAs on mouse chromosomes. J. Struct. Func. Genomics 2 pre, L72-L86 (2001)
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Please visit our web site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 further details.
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onno,H., Fukunishi,Y., Shibata,K., Itoh,M., Carninci,P., Sugahara
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prepared and sequenced in Mouse Genome Encyclopedia
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/clone="A430110L22"
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-DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15 -MODE=LOCAL
-OUTEMT=pto -NORM=ext -HEAPSIZE=500 -MILEN=0 -MAXLEN=20000000000
-USER=US09993234 @CGN_1_1_8534_@runat_27032003_115455_15349 -NCPU=6 -ICPU=3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120
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Yu,G.-L., Ni,J., Gentz,R.L. and Dillon,P.
Death domain containing receptors
Patent: US 6153402-A 3 28-NOV-2000;
Location/Qualifiers
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                                                                                                                                                               PheHisLysLysIleGlyLeuPheCysCysArgGlyCysProAlaGlyHisTyrLeuLys
                                      AlaSerGlnValAlaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLys
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Signal transduction by DR3, a death domain-containing receptor related to TNFR-1 and CD95
Science 274 (5289), 990-992 (1996)
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Chinnaiyan, A.M., O'Rourke, K.,
Duan, R., Xing, L., Gentz, R., N
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Direct Submission
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/product="death receptor 3"
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TLTYTYRHCWPHKFLVTADEAGMEALTPPPATHLSPLDSAHTLLAPPDSSEKICTVQL
VGNSWTPGVPETQEALCPQVTWSWDQLPSRAIGPAAAPTLSPESPAGSPAMMIQPGPQ
LYDVMDAVPARRWKEFVRTLGLREAEIEAVEVEIGRFRDQQYEMLKRWRQQQPAGLGA
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Degli-Esposti,M.A. and Googwin,m.c..
Direct Submission
Submitted (12-NOV-1996) Biochemistry, I
Submitted (12-NOV-1996) WA 98101, USA
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Degli-Esposti, M.A., Din, W.S., Cosman, D., Smith, C.A.
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Location/Qualifiers
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1 (bases 1 to 1254)
Screaton,G.R., Xu,X.N., Olsen,A.L., Cowper,A.E., Tan,R., McMichael,A.J. and Bell,J.I.

LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing
Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
                                         9DU,
                                                          Direct Submission
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                                            GlnMetPheTrpValGln 186
                                                                                                                       GATACTGACTGTGGGACCTGCCTGCCTGGCTTCTATGAACATGGCGATGGCTGCGTGTCC
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ALENCSAVADTRCGCKPGWEVECQVSQCVSSSPFYCQPCLDCGALHRHTRLLCSRRDT
DCGTCLDGFYEBGDGCVSCPTSTLGSCDERCAAVCGWRQMFWQVLLAGLVWPLLLGA
TLTYTYRHCWPHKPLVTADEAGMEALTPPPATHLSPLDSAHTLLAPPDSSEKICTVQL
VGNSWTPGYPETQEALCPQVTMSWDQLLSRALGPAAAPTLSPESPAGSPAMHLQPGPG
LYDVMDAVPARRWKEFVRTLGLREAEIEAVEVEIGRFRDQQYEMLKRWRQQQPAGLGA
VYAALERMGLDGCVEDLRSRLLQRGP"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /function="mediates apoptosis"
/note="LARD-la; membrane protein;
/ns-R1; contains a death domain"
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/protein_id="AAC51306.1"
/db_xref="GI:2071949"
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/cell_type="lymph
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Submitted (19-MAR-1997) Molecular Immunology Group, Institute of
Molecular Medicine, John Radcliffe Hospital, Headington, Oxford
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Screaton,G.R., Xu,X.N., Olsen,A.L., Cowper,A.E., Tan,R., McMichael,A.J. and Bell,J.I.
LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
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Human lymphocyte associated receptor of death
alternatively spliced, complete cds
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Screaton, G.R.
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/protein_id="AcC51307.1"
/db_xref="Gi:2071951"
/translation="MEQRPRGCAAVAAALLLVLLGARAQGGTRSPRCDCAGDEHKKIG
LFCCRGCRGFUKAPCTEGGRSTCLVCPQDTFLAWENHHUSECARCQACDEQASQV
ALENCSAVADTROGCKPGWFUEQVSQCVSSSPYCQPCLDGGALHRHTRLLCSRRDT
DCGTCLPGFYEHGDGCVSCPTSTLGSCPERCAAVCGWROMFWVOYLLAGLVVPLLLGA
TLTYTYRHGWPHKPLVTAADEAGMEALTPPPA7HLSPLDSAFTLLAPPDSSEKICTVQ
LVGNSWTPGYPETQGALCPQVTWSWDQLPSRALGPAAAPTLSPESPAGSPAMMLQPGP
QLYDVMDAVPARRWKEFVRTLGLREAETEAVEVEIGRFRDQQYEMLKRWRQQQPAGLG
AVYAALERMGLDGCVEDLRSRLQRGP"
                                                                                                                                                                                                                                                                                                                                                                                                     /note="insertion compared to LARD-1a, deposited in GenBank Accession Number U94501, probably represents alternative 3' splice site"
1 421 c 408 g 226 t
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TNF-R1; contains a death domain"
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/db_xref="taxon:9606"
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U74611
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Marsters (S.A., Sheridan, J.P., Donahue, C.J., Pitti, R.M., Gr
Goddard, A.D., Bauer, K.D. and Ashkenazi, A.
Apo-3, a new member of the tumor necrosis factor receptor
contains a death domain and activates apoptosis and NF-kB
                                                                                                                                                                                                          Submitted (15-OCT-1996) Molecular Bruno Blvd., South San Francisco, Location/Qualifiers
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          /product="Apo-3"
/product="Apo-3"
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DCGTCLPGFYEHGDGCVSCPTSTLGSCPERCAAVCGWRQMFWVQVLLAGLVVPLLLGA
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/dev_stage="fetal"
89. .1342
                                                                                               /note="contains death domain, NF-kB; TNF receptor family mer
                                                                                                                                                                /organism="Homo sapiens"
/db_xref="taxon:9606"
/chromosome="1"
                                                                                                                                                        /map="1p36.3"
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                                                                                                                                                                                                                                                                                                                                                      ; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                          Homo sapiens
Eukaryota; Metazoa; C
Mammalia; Eutheria; P
Mammalia; Lto 1662)
                                                                                                                                                        Sequence
AX055442
Ashkenazi, A.J., Baker, K.P., Chan, B., Gurney, A.L., Hebert, C., Henzel, W., Katumas, D., Watanabe, C.K. and Wood, W.I.
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VYAALERWGLDGCVEDLRSRLORGP
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Matches:
Conservative:
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                                                             Craniata; Vertebrata; Catarrhini; Hominidae;
                Kabakoff,R.C.,
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                Godowski, P.J., Shelton, D.L.,
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Mammalia; Eutheria; Primates; Ca

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Ashkenazi, A.J., Goddard, A., God
Hillan, K.J., Marsters, S.A., Pan
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Indels:
Gaps:
   Godowski,P.J., Gurney,A.L., Pan,J., Pitti,R.M., Roy,M.A.,
                       Craniata; Vertebrata; Catarrhini; Hominidae;
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Mammalia; Eutheria; Primates;
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/protein_id="AAB41432.1"
/protein_id="AAB41432.1"
/db_xref="GI:1800293"
/translation="GCAAVAAALLLVLLGARAQGGTRSPRCDCAGDFHKKIGLFCCRG
CPAGHYLKAPCTEPCGNSTCLVCPQDTFLAWENHHNSECARCQACDEQASQVALENCS
AVADTRCGCKFGWFYECQVSGCVSSSPFYCQPCLDCGALHRHFTLLCSRRDTPCGTCL
LGFYEHGDGCVSCPTSTLGSCPERCAAVCGWRQMFWVQVLAGLVVPLLLGATLTYTY
RHCWPHKPLVTADEAGMEALTPPPATHLSPLDSAHTLLAPDDSSEKICTVQLVGNSWT
PGYPETQEALCPQVTWSWDQLPSRALGPAAAPTLSPESPAGSPAMMLQPGPQLYDVMD
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/function="mediates apoptosis when cross-linked"
/function="mediates apoptosis when cross-linked"
/note="A TER1-related death-domain containing receptor;
/note="A TER1-related death-domain containing receptor;
DR3, WSL-LR, Apo-3, TRAMP; transmembrane form; similar to
human EST clone 298913, GenBank Accession Number N71143"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /gene="DDR3"
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/map="1p36"
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Bodmer, J.L., Burns, K., Schneider, P., Hofmann, K., Steiner, V.,
Thome, M., Bornand, T., Hahne, M., Schroter, M., Wilson, A.,
French, L.E., Browning, J.L., MacDonald, R. and Tschopp, J.

Direct Submission
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Bodmer, J. L., Burnas, K., Schneider, P., Hofmann, K., Steiner, V., French, L. E., Browning, J. L., MacDonald, R. and Tschopp, J. TRAMP, a novel apoptosis-mediating receptor with sequence homology to tumor necrosis factor receptor 1 and Fas(Apo-1/CD95)

Immunity 6 (1), 79-88 (1997)
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                                                                                                                                                                                 /translation="AlagCaavaaalllvilgaraoggtrsprcdcagdffkkiglfc
CRGCpaghtikapctepcgnstcivcpodtflawenhhnsecarcoacdegasqvale
NCSAVADTRCGCKPGWFVECOVSQCVSSSPFYCOPCLDCGALHRITALCSRRDTDCG
TCLLGFYEREDGCVSCPTSTLGSCPERCAAVCGWRQMFWVQVLLAGLVVPLLLGATIDCG
YTYRHCWPHKPLVTADEAGMEAITPPPATHLSPLDSAHTLLAPDDSSEKICTVQLUGN
SWTGYSPETQEALCPQVTWSWDQLPSRALGPAAAPTLSPESPAGSPAMMLOPGPQLYGN
VMDAVPARRWKEFVRTLGLREAEIEAVEVEIGRFRDQQYEMLKRWRQQQPAGLGAVYA
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                                                                                                                                                                                                                                                                                       /product="apoptosis-mediating receptor TRAMP"
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/db_xref="GI:1695925"
                                                                                                                                                                                                                                                                                                                                     /note="contains a death domain; similar to TNF receptor"
                                                                                                                                                                                                                                                                                                                                                           function="mediates apoptosis"
                                                                      1.29e-67
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                                                                                                                                                                                                                                                                                                                                                                                                                                 /chromosome="1"
                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ocation/Qualifiers
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181 GlnMetPheTrpValGln 186
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                                                                                                                                      Direct Submission
Submitted (19-MAR-1997) Molecular Medicine, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Screaton,G.R., Xu,X.N., Olsen,A.L., Cowper,A.E., Tan,R., McMichael,A.J. and Bell,J.I.
LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing
Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae;
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                                               /db_xref="taxon:9606"
/cell_line="HeLa"
                                                                                                            Location/Qualifiers
                                                                             'organism="Homo sapiens"
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DCGTCLPGFYEHGDGCVSCPTSTLGSCPERCAAVCGWRQSRWCAGNASGRTGMDRGEA
GEEGGNHPTPTSCFQCSGSRCSWLALWSPSCLGPP"
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            2456
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Accession Number U94501, probable retained intron leading
to premature transcriptional termination"
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Horrigan, S., Soppet, D.R. and Weaver, Z.
Cancer gene determination and therapeutic
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Submitted (12-NOV-1996) J. Kitson, GLAXO-Wellcome,
Research Centre, Gunnels Wood Road, Stevenage, SGI
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TLTYTYRHCWPHKFLYTADEAGMEALTPPATHLSPLDSAHTLLAPDSSEKICTYOLLLGA
VONSWTPCYPETOGALCPQVTWSWDQLPSRALGPARAPTLSPESPAGSPAMMLQPGPQ
LYDVMDAVPARRWKEFVRTLGLREAEIEAVEVEIGLFRDQQYEMLKHWRQQQPAGLGA
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/protein_id="CAA70560.1"
/db_xref="GI:1669692"
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/protein_id="CAA70561.1"
/db_xref="GI:1669693"
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/tissue_type="lymphoid"
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/db_xref="taxon:9606"
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             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 1143)
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                                                                                                                          TTGGCCTGGGAGAACCACCATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAG
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Submitted (19-MAR-1997) Molecular Immunology Group, Institute of
Molecular Medicine, John Radcliffe Hospital, Headington, Oxford
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LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note="deletion compared to LARD la, deposited in GenBank
Accession Number U94501, probable skipping of putative
exon 7 encoding the transmembrane domain"
378 c 378 g 199 t
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DSAHTLLAPPDSSEKICTVQLVGNSWTPGYPETQEALCPQVTWSWDQLPSRALGPAAA
PTLSPESPAGSPAMMLQPGPQLYDVMDAVPARRWKEFVRTLGLREAEIEAVEVEIGRF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /translation="MEORPRGCAAVAAALLLVLLGARAOGGTRSPRCDCAGDFHKKIG
LFCCRGCPAGHYLKAPCTEPCGNSTCLVCPQDTFLAWENHHNSECARCQACDEQASOV
ALENCSAVADTRCGCKPGWFVECQVSQCVSSSPFYCQPCLDCGALHRHTRLLCSRRDT
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/protein_id="AAC51315.1"
/db_xref="GI:2071967"
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/note="LARD-9; NGFR family member;
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/cell_line="HeLa"
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                                                                                                   TGCCCCACGAGCACCCTGGGGAGCTGTCCAGAGCGCTGTGCCGCTGTCTGGCCTGGAGG
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SUMMARIES

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GenCore version Copyright (c) 1993 - 2003
5.1.4_p5_4578
Compugen Ltd.
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Command line parameters:

-MODEL_frame+_D2n.model -DEV=xlh
-Q-cgn2_1/USPTQ_Spool/US09993234/runat_27032003_115454_15312/app_query.fasta_1.2346
-Q-ycgn2_1/USPTQ_Spool/US09993234/runat_27032003_115454_15312/app_query.fasta_1.2346
-DB=N.Geneseq_101002 -QFMT=fastap -SUFFIX=rng -MINKATCH=0.1 -LOOPCL=0
-LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi
-LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15
-MODE-LOCAL -OUTEMT=pto -NORM-ext -HEADSIZE=500 -MINLEN=0 -MAXLEON-000000000
-USER=US09993234 @CGN 1_1_660_@runat_27032003_115454_15312 -NCPU=6 -ICPU=3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPDEXT=0.5 -FGAPOP=6 -FGAPEXT=7
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Homo sapiens

Death domain containing receptor; DR3; human; apoptosis; inflammation; NF-kappaB; ds.

RESULT 1
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          101
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P-PSDB; AAW31517.
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ProGlyTrpPheValGluCysGlnValSerGlnCysValSerSerSerProPheTyrCys 120
                        LeuLeuGlyAlaArgAlaGlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAsp
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NF-kappaB, antagonists can be
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           The invention provides nucleotide sequences encoding death domain containing receptor polypeptides DR3-V1 and DR3. The DR3-V1 cDNA clone is contained in ATCC deposition No. 97456 and the DR3 cDNA clone is contained in ATCC deposition No. 9757. Recombinant vectors comprising the nucleic acid sequences and optionally the leader sequences are used for the recombinant production of the proteins. The present sequence represents a cDNA encoding the death domain containing receptor polypeptide (DR3).
                                                                                                                                                                                                                    06-FEB-1997;
12-MAR-1996;
17-OCT-1996;
                                                                                                        Claim 6; Fig 3; 50pp; Japanese.
                                                                                                                          New death domain containing receptor and optionally comprising leader sequence
                                                                                                                                                                    WPI; 1999-124390/11.
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96US-0013285.
96US-0028711.
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Sequence

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201 A; 420

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RESULT 3
AAC6877
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                       21-APR-2000;
                                                                                                                                                                                                   neurodegenerative
                                                                                                                                                                                                                           autoimmune
                                                                                                                                                                                                                                                Human; death
                                                                                                                                                                                                                                                                                          Human death
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                                                                                                                                                                                                                           disorder;
                                                                                                                                                                                                                                                                                       domain
                                                                                                                                                                                                                                           domain containing receptor; DR3; cancer;
                       2000WO-US10741
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                                                                                                                                                                                                   disease;
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The present invention provides the protein and coding sequences for two death domain containing receptors, designated DR3 and DR3-VI. These receptors are involved in apoptosis, and the sequences given can be used in the treatment of cancers, infections, cardiovascular disorders such as arrhythmias, ischaemia, aneurysms, arterial occlusive diseases, embolisms and congenital heart defects, neurodegenerative diseases including Alzhelmer's and Parkinson's diseases, autoimmune disease such as multiple sclerosis, arthritis, diabetes, Graves' disease, asthma and psoriasis, and to promote angiogenesis and wound healing.
                                                                                                                                                                         Treating graft-versus-host disease, cancer, immunodeficiency autoimmune disease comprising administering an antibody to De Containing Receptor proteins and a second therapeutic agent .
Sequence 1254 BP; 201 A; 420
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                                                                                                                                                    Example 2B;
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28-MAY-1999;
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DB; AAB36265.
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99US-0136741.
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Domain

Best Local Similarity: Query Match:
DB: US-09-993-234-6\_COPY\_19\_204 (1-186) Score: Alignment Scores: Percent Similarity: М О.: 7.95e-75 1101.00 100.00% 100.00% 100.00% 21 x AAC68777 (1-1254) Length:
Matches:
Conservative:
Mismatches:
Indels:

₽ Ş 밁 QV 115 21 PheHisLysLysIleGlyLeuPheCysCysArgGlyCysProAlaGlyHisTyrLeuLys 55 1 LeuLeuGlyAlaArgAlaGlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAsp CTGCTGGGGGCCCGGGCCCAGGCGCACTCGTAGCCCCAGGTGTGACTGTGCCGGTGAC TTCCACAAGAAGATTGGTCTGTTTGTTGCAGAGGCTGCCCAGCGGGGCACTACCTGAAG 40 174 114

175 41 AlaProCysThrGluProCysGlyAsnSerThrCysLeuValCysProGlnAspThrPhe GCCCCTTGCACGGAGCCCTGCGGCAACTCCACCTGCCTTGTGTGTCCCCCAAGACACCTTC

235 61 LeuAlaTrpGluAsnHisHisAsnSerGluCysAlaArgCysGlnAlaCysAspGluGln TTGGCCTGGGAGAACCACCATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAG

294 80 234 60

8. B

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101 414 120

140 474

AspThrAspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSer

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C CDNA clone FH20.57 (ATCC 55820) codes for a novel polypeptide (see 20 w26709), designated Apo-3, that stimulates or induces apoptotic 2 activity in mammalian cells. It was isolated from a human foetal 3C heart cDNA library by screening with probes (see T91183-84) based 2C on an EST sequence (GenBank locus W71984) that showed homology to 2C the intracellular domain of human TNFRI and CD95. Amino acid 3C con be used diagnostically for tissue-specific typing and to 2C polypeptide, Apo-2LI (see W26708). Nucleic acids encoding Apo-3 3C can be used diagnostically for tissue-specific typing and to 2C produce recombinant Apo-3 polypeptides, especially the 2C extracellular domain (amino acids 1-198) or death domain (amino 38-417). Apo-3 can be used to induce apoptosis or 38-417). Apo-3 can be used to induce apoptosis or NF-Kappa-N- or JNK-mediated gene expression for therapeutic 2C purposes. Non-human transgenic animals containing cells that acid, and knockout animals containing 3C cells that have an altered Apo-3 gene, can be used in drug 3C screening and development.
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AAT9118
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                                                                                                                                                                                                                                                                                                                                        Ashkenazi
                                                                                                                                                                                                                                                                                                                                                                                      23-SEP-1996;
01-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                            31-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                              (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-OCT-1997.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       apoptosis;
                                                                                                                                                                                                                                                  Page 45-46;
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96US-0625328
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89..160
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/note= "determined by hydropathy analysis"
161.1339
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89..1342
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Sequence

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                                                                                          Rheumatoid arthritis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI;
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                                                                                                                                                                                                                                                                                                                       LeuLeuGlyAlaArgAlaGlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAsp
                                               AspThrAspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSer
                                                                                      GlnProCysLeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArg
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                                      GATACTGACTGTGGGACCTGCCTGCCTGGCTTCTATGAACATGGCGATGGCTGCGTGTCC
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27-MAR-2001;
30-MAR-2001;
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                                                                                                                                                                                                                                                                      Genomes,
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SHIOZAWA S.
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2001JP-0099990.
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29-OCT-1999
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2000WO-US07532.

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2000WO-US13705.
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Ashkenazi Hebert C, Wood WI; ΑJ, NJ, Baker KP, Chan B, Henzel W, Kabakoff RC, Goddard A, (Shelton DL, Godowski PJ, , Tumas D, V Watanabe

WPI; 2001-025253/03. P-PSDB; AAB50918.

Thirty three nucleic acids encoding PRO polypeptides which are useful in the diagnosis and treatment of immune related disorders, e.g. systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis, thyroiditis and diabetes mellitus

Claim 48; Fig 33; 218pp; English.

The present sequence is one of thirty three nucleic acids encoding PRO CC polypeptides. The PRO polypeptides, anti-PRO antibodies, agonists and cantagonists are useful for treating and diagnosing immune related CC disorders such as systemic lupus erythematosus, rheumatoid arthritis, costeoarthritis, juvenile chronic arthritis, spondyloarthropathies, systemic vasculitis, autoimmune haemolytic costeoarthropathies, idiopathic inflammatory myopathies, Sjogren's cc syndrome, systemic vasculitis, sarcoidosis, autoimmune haemolytic commune-mediated renal disease, demyelinating diseases of the central cc immune-mediated renal disease, demyelinating diseases of the central cc immune-mediated renal disease, demyelinating diseases and chronic cc inflammatory demyelinating polyneuropathy or Guillain-Barre syndrome, and chronic cl ciscome, and chronic active hepatitis, primary cirhosis, granulomatous hepatitis and sclerosing cholangitis), cc disease, autoimmune or immune-mediated skin diseases (such as bullous skin diseases, erythema multiforme, contact dermatitis, proriasis), cc disease, autoimmune or immune-mediated skin diseases (such as bullous skin diseases, erythema multiforme, contact dermatitis, proriasis), cf food hypersensitivity and urticaria), immunological diseases of the cc and hypersensitivity pneumonits), transplantation associated diseases cc including graft rejection and graft-versus-host diseases.

1662 B₽; 302 A; 540 C; 531 G; 289 T; 0 other;

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41 AlaProCysThrGluProCysGlyAsnSerThrCysLeuValCysProGlnAspThrPhe 	21 PheHisLysLysIleGlyLeuPheCysCysArgGlyCysProAlaGlyHisTyrLeuLys	1 LeuLeuGlyAlaArgAlaGlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAsp 	US-09-993-234-6_COPY_19_204 (1-186) x AAC91477 (1-1662)	pred. No.: Pred. No.: Score: Percent Similarity: Best Local Similarity: Query Match: DB:
LuProCysGlyAsnSe 	LeGlyLeuPheCysCy               TGGTCTGTTTTGTTC	rgAlaGlnGlyGlyT]              3GGCCCAGGGCGGCAC	_204 (1-186) x /	1.1e-74 1101.00 100.00% 100.00% 100.00% 22
PTThrCysLeuValCy	/SArgGlyCysProAl              CAGAGGCTGCCCAGC	hrärgSerProärgCy               TCGTAGCCCCAGGTC	AAC91477 (1-166;	Length: Matches: Conservative: Mismatches: Indels: Gaps:
AlaProCysThrGluProCysGlyAsnSerThrCysLeuValCysProGlnAspThrPhe	PheHisLysLysIleGlyLeuPheCysCysArgGlyCysProAlaGlyHisTyrLeuLys 	LeuLeuGlyAlaArgAlaGlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAsp 	2)	1662 186 0 0 0

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US-09-993-234-6_COPY_19_204 (1-186) x AAT89426 (1-1783)
                                                                              Alignment Scores: Pred. No.: Score:
                               Query Match:
DB:
                                                  Percent Similarity:
Best Local Similarity:
                                                                                                                                             This cDNA clone, deposited as ATCC 97456, codes for human death domain containing receptor DR3-V1 (see AAW31516), a novel member of the tumour necrosis factor receptor family. It was isolated from a CCDNA library derived from cells of a human testis tumour. Related containing receptor DR3 cDNA (see AAR8427) was isolated from a HUVEC cDNA library. The genes have also been identified in CC CDNA libraries of foetal liver, foetal brain, tonsil and leukocyte. CCNA libraries of foetal liver, foetal brain, tonsil and leukocyte. CC nucleic acids encoding full-length or mature DR3-V1 can be used to produce recombinant polypeptides in transformed host cells. These CC polypeptides can be used to treat diseases and disorders associated with the inhibition of apoptosis. Antagonists, such as antibodies craised against DR3-V1, can be used to treat diseases and disorders cc associated with increased apoptosis and for treating inflammatory could be considered with increased apoptosis and for treating inflammatory could be considered with increased apoptosis and for treating inflammatory could be considered.
                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                              Claim
                                                                                                                                                                                                                                                                                                                                                             Death domain containing receptor polypeptide(s) DR3 and DR3-V1 for activation of apoptosis and NF-kappaB, antagonists can be to treat inflammatory diseases
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97US-0037341.
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RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention provides nucleotide sequences encoding death domain containing receptor polypeptides DR3-VI and DR3. The DR3-VI cDNA clone is contained in ATCC deposition No. 97456 and the DR3 cDNA clone is contained in ATCC deposition No. 97757. Recombinant vectors comprising the nucleic acid sequences and optionally the leader sequences are used for the recombinant production of the proteins. The present sequence represents a cDNA encoding the death domain containing receptor polypeptide (DR3-VI).
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                                                                                                               1999-124390/11.
DB; AAW95537.
                                   GlnMetPheTrpValGln
                                                                  GlnProCysLeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArg
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                                                                                                                                                                                                        US-09-993-234-6_COPY_19_204 (1-186)
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                                                                                                                                                                                                                                                                                                                                                                                                                          Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention provides the protein and coding sequences for two death domain containing receptors, designated DR3 and DR3-V1. These receptors are involved in apoptosis, and the sequences given can be used in the treatment of cancers, infections, cardiovascular disorders such as arrhythmias, ischaemia, aneurysma, arterial occlusive diseases, embolisms and congenital heart defects, neurodegenerative diseases including alzheimer's and Parkinson's diseases, autoimmune disease such as multiple sclerosis, arthritis, diabetes, Graves' disease, asthma and psoriasis, and teachers and teachers of the sequence of the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; death domain containing recautoimmune disorder; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Treating graft-versus-host disease, cancer, immunodeficiency autoimmune disease comprising administering an antibody to De Containing Receptor proteins and a second therapeutic agent .
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28-MAY-1999;
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DB; AAB36264.
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RESULT 12
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     Claim 2;
              DNA encoding apoptosis inducing receptor - which is Type transmembrane protein, useful for regulating cell death
                                      WPI; 1998-240077/21.
P-PSDB; AAW57045.
                                                           Perkins PA;
                                                                                     04-OCT-1996;
                                                                                                   03-OCT-1997;
                                                                        (IMMV ) IMMUNEX
                                                                                                                  09-APR-1998
                                                                                                                              W09814565-A1
                                                                                                                                                                                               Apoptosis inducing receptor; AIR protein; Type I transmembrane protein; tumour cell
                                                                                                                                                                                                                 Human apoptosis inducing receptor coding sequence.
                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                           therapy; ss.
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     Page 28-30;
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                                                                                                  97WO-US17876
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/product= AIR
                                                                                                                                                         Location/Qualifiers 236..1489
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  45pp; English
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death;
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RESULT 13
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                         Breast
                                               15-MAY-2002
                                                                                             ABL64119 standard; DNA;
                                                                        ABL64119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This sequence encodes the human apoptosis inducing receptor (AIR) of the invention. AIR is a Type I transmembrane protein, soluble forms of which can be used to regulate cell death in a therapeutic setting. Soluble AIR can also be used in vitro to block apoptosis or AIR-expressing cells, or to screen agonists or antagonists of AIR activity. The cytoplasmic domain of AIR can be used to develop assays for inhibitors of AIR-induced cell death, which is useful to regulate cell death in a therapeutic setting as well as in vitro. Agonists of AIR activity can be used to kill tumour cells that express AIR, or T cells expressing AIR in autoimmune diseases.
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                                                                                                                                                                                                                                                    AspThrAspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSer
                                                                                                                                                                                                                                                                                                    GlnProCysLeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArg
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Matches:
Conservative:
Mismatches:
Indels:
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                   SEQ ID NO:2456.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              347 T; 0 other;
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186
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Human;

cancer;

colon; breast;

ovary; oesophagus; kidney; thyroid;

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The present inventic anti-neoplastic agen agent to be tested f expression of at leacomprises a sequence
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18-SEP-2000;

18-SEP-2000;

20-SEP-2000;

20-SEP-2000;

21-SEP-2000;

22-SEP-2000;

25-SEP-2000;

25-SEP-2000;

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20-CCT-2000;

20-CCT-2000;

21-CCT-2000;

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21-SEP-2000;

22-SEP-2000;

23-SEP-2000;

24-SEP-2000;

25-SEP-2000;

26-SEP-2000;

27-SEP-2000;

28-SEP-2000;

29-SEP-2000;

29-SEP-2000;

20-CCT-2000;

20-CCT-20
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cytostatic; gene therapy;
gene; ds.
                                                                                                                                                                                                                                                           Screening for anti-neoplastic agent involves exposing cells to a chemical agent to be tested for anti-neoplastic activity, and determining a change in expression of a gene of a signature gene
                                                                                                                                                                                                      Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Young
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Weaver z
invention describes a method (M1) for screening f
stic agent. The method involves exposing cells to
tested for anti-neoplastic activity, determining
of at least one gene (I) of a signature gene set,
sequence (S) selected from 8447 sequences (given
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2000US - 231133P.
2000US - 233617P.
2000US - 234034P.
2000US - 234567P.
2000US - 23563P.
2000US - 23663P.
2000US - 23603P.
2000US - 23603P.
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2000US - 23603P.
2000US - 237609P.
2000US - 237713P.
2000US - 237716P.
2000US - 237716P.
2000US - 237716P.
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; antineoplastic; Wilm's tumour; adenocarcinoma;
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         for an control of cont
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         RESULT 14
AAF83770
ID AAF83
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AC AAF83
XX
DT 06-AU
XX
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to ABL70110), or is at least 95% identical to (5), where a change in expression is indicative of anti-neoplastic activity. (I) has cytostatic activity and can be used in gene therapy. MI can be used for screening an anti-neoplastic agent, and can be used for producing a product which is the data collected with respect to the anti-neoplastic agent as a result of MI, and the data is sufficient to convey the chemical structure and/or properties of the agent. MI can be used in the treatment of cancer such as colon, breast, stomach, lung, thyroid, oesophageal, ovarian, Kidney, prostate or pancreatic cancer, infiltrating lobular cancer, squamous cell carcinoma, neuroendocrine carcinoma, papillary carcinoma and Wilm's tumour.

Sequence 1743 BP; 347 A; 543 C; 549 G; 304 T; 0 other;
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Alignment Scores:
Pred. No.:
Score:
Percent Similarity:
Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1743
AAF83770
                                                                                                                                                                                                                                                                                                                                                                                                        1 LeuLeuGlyAlaArgAlaGlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAsp
                                                                            CysProThrSerThrLeuGlySerCysProGluArgCysAlaAlaValCysGlyTrpArg
                                                                                                                                                        AlaSerGlnValAlaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLys
                                                                                                                                                                                                                                                                            GlnMetPheTrp
                                                                                                                            AspThrAspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSer
                                                                                                                                                                                                          ProGlyTrpPheValGluCysGlnValSerGlnCysValSerSerSerProPheTyrCys
                                       CAGAGTAGGTGG
                                                                                                                   GATACTGACTGTGGGACCTGCCTGCCTTGCTATGAACATGGCGATGGCTGCGTGTCC
                                                                                                                                                                                                CCAGGCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGC
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98.91%
97.82%
24
DNA;
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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(first entry)

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Query Match:
               Best Local
                          Percent Similarity:
                                                       Pred.
                                                                     Alignment Scores
                                                                                                                                   The invention relates to a composition comprising a biologically active TR3-specific binding agent (I) that binds to TR3 and inhibits the proliferation of cells expressing TR3. (I) identified by the methods are useful for treating a subject suspected of having a disease associated with a proliferation of cells expressing TR3 especially leukemias or lymphomas or a T-cell mediated disease especially autoimmune diseases such as myasthenia gravis, systemic lupus erythamatosus, rheumatoid arthritis, diabetes, multiple sclerosis, sarcoidosis, myocarditis, thyroiditis and tumours. (I) is also useful for treating a subject suspected of having graft-versus-host disease, rejection of a transplanted organ such as heart, liver, lung, kidney, pancreas, bowel, skin or an appendage, or inflammatory diseases, allergies and contact dermatitis. The present sequence represents the nucleotide sequence of
                                                                                                                                                                                                                                                                                                                                                      Composition for treatment of T-cell mediated disease e.g. arthritis, cancer comprises a biologically active TR3-specific binding agent especially a monoclonal antibody - \frac{1}{2}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo
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                                                                                                                                                                                                                                                                                                                                                                                                             P-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleotide sequence
                                                                                                                                                                                                                                                                                                                            Disclosure; Page 72;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-NOV-1999;
                                                       No . :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sapiens
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               Similarity:
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antiinflammatory;
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/transl_except= "(pos:558..559, aa:Cys)"
/note= "this codon has an apparent 1 nucleotide
deletion which alters the reading frame"
/transl_except= "(pos:638..639, aa:Leu)"
/note= "this codon has an apparent 1 nucleotide
deletion which alters the reading frame"
/transl_except= "(pos:718..719, aa:Met)"
/note= "this codon has an apparent 1 nucleotide
deletion which alters the reading frame"
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73..1247
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1...1250
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antidiabetic; neuroprotective; cardiant;
antiallergic; T-cell-inhibitor; ds.
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05-JUN-2000;
05-JUN-2000;
18-SEP-2000;
18-SEP-2000;
20-SEP-2000;
20-SEP-2000;
                                                                                                                                             cytostatic;
                                                                                                                                                     stomach;
                                                                                                                                                                                 Thyroid cancer
                                                                                                                                                                                                                   ABL67258;
                                                               30-MAY-2001;
                                                                                 13-DEC-2001
                                                                                                  WO200194629-A2
                                                                                                                   Homo sapiens.
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                                                                                                                                                      cancer;
                                                                                                                                            er; colon; breast; ovary; oesophagus; kidney; thyroid; ng; prostate; pancreas; carcinoma; antitumour; cancero gene therapy; antineoplastic; Wilm's tumour; adenocar
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2000US-209531P.
2000US-233133P.
2000US-233617P.
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                                                                                                                                                                               related gene sequence SEQ ID NO:5595.
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The present invention describes a method (MI) for screening for an CC anti-neoplastic agent. The method involves exposing cells to a chemical capent to be tested for anti-neoplastic activity, determining a change in CC expression of at least one gene (I) of a signature gene set, where (I) CC comprises a sequence (S) selected from 8447 sequences (given in ABL61664 CC to ABL70110), or is at least 95% identical to (S), where a change in CC expression is indicative of anti-neoplastic activity. (I) has cytostatic CC activity and can be used in gene therapy MI can be used for screening CC an anti-neoplastic agent, and can be used for producing a product which is the data collected with respect to the anti-neoplastic agent as a CC result of MI, and the data is sufficient to convey the chemical CC structure and/or properties of the agent. MI can be used in the CC treatment of cancer such as colon, breast, stomach, lung, thyroid, CC oesophageal, ovarian, kidney, prostate or pancreatic cancer, cfilitrating ductal cancer, clast cell cancer, infiltrating ductal cancer, CC infiltrating lobular cancer, squamous cell carcinoma, neuroendocrine CC carcinoma, papillary carcinoma and Wilm's tumour.
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Best Local Similarity:
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-DB=ISSued_Patents_NA -QFMT=fastap -SUFFIX=rn1 -MINMATCH=0.1 -LOOPCL=0
-LOOPEXT=0 -UNITS-bits -START=1 -END=-1 -MATRIX-blosum62 -TRANS-human40.cdi
-LIST=45 -DCALIGN=200 -THR_SCORE-pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15
-MODE-LOCAL -QUTFMT=pto -NONH-ext -HEAPSIZE-500 -MINLEN=0 -MAXLEN=200000000
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US-08-928-069-2

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### ALIGNMENTS

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                                                                    APPLICATION NUMBER: US/08/
FILING DATE: HEREWITH
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: No. 61
FILING DATE: 06-FEB-1997
PRIOR APPLICATION DATA:
                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Dixit, Vishva
APPLICANT: Gentz, Reiner L.
APPLICANT: Dillon, Patrick J.
TITLE OF INVENTION: Death Dom
                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
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                                   APPLICATION NUMBER: FILING DATE: 17-OC'
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ZIP: 20005-3934
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RESULT 2
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Sequence
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SEQUENCE CHARACTERISTICS:
LENGTH: 1254 base pairs
TYPE: nucleic acid
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ATTORNEY/AGENT INFORMATION:
NAME: Steffe. Fria "
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LOCATION:
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STRANDEDNESS: doub
TOPOLOGY: both
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REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 14
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                                                              GlnMetPheTrpValGln 186
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                                                                                                              CysProThrSerThrLeuGlySerCysProGluArgCysAlaAlaValCysGlyTrpArg
                                                                                                                                                                 AspThrAspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSer
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Application US/08928069
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DB:
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INFORMATION FOR SEQ ID NO: 11.
SEQUENCE CHARACTERISTICS:
FRIGTH: 1634 base pairs
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PRIOR APPLICATION NUMBER: 60/026943
APPLICATION NUMBER: 60/026943
FILING DATE: 09/23/1996
ATTORNEY,AGENT INFORMATION:
NAME: MATSCHARP, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P105
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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CURRENT APPLICATION DATA:
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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STRANDEDNESS: Sing
TOPOLOGY: Linear
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CITY: South San F:
STATE: California
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ProGlyTrpPheValGluCysGlnValSerGlnCysValSerSerSerProPheTyrCys 120
                                                          AlaSerGlnValAlaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLys 100
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RESULT 3
US-08-828-683A-9
US-08-828-683A-9
Sequence 9, Application US/08828683A
Patent NO. 6469144
Patent NO. 6469144
GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
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SEQUENCE CHARACTERISTICS:
LENGTH: 1634 base pairs
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FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
ATTORNEY/AGENT INFORMATION:
NAME: MAISCHAMP, Diane L.
REGISTRATION NUMBER: 35,600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: U5/08/828,683A FILING DATE: 31-Mar-1997 CLASSIFICATION: <UNknown> PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
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TELEPHONE: 650/225-5416
TELEFAX: 650/952-9881
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LeuLeuGlyAlaArgAlaGlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAsp
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STRANDEDNESS: Single
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, VG

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/815,469
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                      APPLICATION NUMBER: NO. FILING DATE: 06-FEB-1997 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Gentz, Reiner L.
APPLICANT: Dillon, Pattick J.
TITLE OF INVENTION: Death Domain
NUMBER OF SEQUENCES: 17
                                                                                                                                                                             CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
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                            APPLICATION NUMBER: US 6 FILING DATE: 17-OCT-1996
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CITY: Washington
STATE: DC
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1100 New York Ave.,
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Gentz, Reiner L.
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Ni, Jian
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APPLICATION NUMBER: US 6 FILING DATE: 12-MAR-1996

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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 141
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
TELEPAX: 202-371-2540
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LOCATION:
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STRANDEDNESS: double
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                                          GlnMetPheTrpValGln 186
                                                                        CysProThrSerThrLeuGlySerCysProGluArgCysAlaAlaValCysGlyTrpArg
                                 CAGATGTTCTGGGTCCAG
                                                                                                                   ProGlyTrpPheValGluCysGlnValSerGlnCysValSerSerSerFroPheTyrCys
                                                                                                                                                                                                                                                                   AlaSerGlnValAlaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLys
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                                                                                                                                                                                                          CCAGGCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGC
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Gaps:
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Matches:
Conservative:
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INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P10
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
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MEDIUM TYPE: 3.5 inch, 1.44 mb 1
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
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LENGTH: 1438 base pair
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NUMBER OF SEQUENCES: 15
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             ProGlyTrpPheValGluCysGlnValSerGlnCysValSerSerSerProPheTyrCys 120
                                                                     AlaSerGlnValAlaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLys 100
                                                                                                                                                                    CCAGGCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGC
                                                                                                                            LeuAlaTrpGluAsnHisHisAsnSerGluCysAlaArgCysGlnAlaCysAspGluGln
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                                                      GCCTCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGGCTGTGGCTGTAAG
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: Nucleic Acid
NDEDNESS: Single
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N: 435
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                               US-09-993-234-6_COPY_19_204 (1-186) x US-08-828-683A-5 (1-1438)
                                                                 Query Match:
DB:
                                                                                               Best Local Similarity:
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                                                                                                               Percent Similarity:
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Patent No. 6469144
GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
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1 LeuLeuGlyAlaArgAlaGlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAsp
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FILING DATE: 31-Mar-1997
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION: TELEPHONE: 650/225-5416 TELEFAX: 650/952-9881
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy
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ADDRESSEE: Genentech,
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                                                                                                                                                                                                                          LENGTH: 1438 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech)
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                                                                                                                                                                                                                                                                                                                                                                 NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007P1
                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 08/625328
FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
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Matches:
Conservative:
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                                                      FILING DATE: 09/23/1996
ATTORNEY/AGENT INFORMATION:
NAME: MATSCHANG, Diane L.
REGISTRATION NUMBER: 35,600
                           REFERENCE/DOCKET NUMBER: P1 TELECOMMUNICATION INFORMATION:
                                                                                                                                      CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                     SOFTWARE: WinPatin (Ge CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inc
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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                                                                                                                         APPLICATION NUMBER:
                                                                                                                                                                       APPLICATION NUMBER: US/08/928,069 FILING DATE: 11-Sep-1997
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                TELEPHONE:
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Best Local Similarity:
Query Match:
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GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-2 LI AND NUMBER OF SEQUENCES: 28
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ADDRESSEE: Genentech, Inc.
STREET: 1 DNA Way
CITY: South San Francisco
STATE: California
COUNTRY: USA
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TYPE: Nucleic Acid
STRANDEDNESS: Single
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MEDIUM TYPE: 3.5 inch, 1.44 mb 1

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/828,683A
FILING DATE: 31-Mar-1997
                                                                                                                                                                                                                                                                                                            CAGAGAT 426
                                                                                                                                                                                                                                                                                                                                     gArgAsp 141
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                                                               Sequence 10,
Patent No. 5
GENERAL INFORMATION:
APPLICANT: Beutler, Bruce A.
APPLICANT: Bazzoni, Flavia M.
TITLE OF INVENTION: MODIFIED RECEPTORS
TITLE OF INVENTION: SIGNAL
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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 433 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
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APPLICATION NUMBER: 08/625328
FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-5ep-1996
ATTORNEY/AGENT INFORMATION:
NAME: Marschang, Diane L.
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REFERENCE/DOCKET NUMBER: P1007P1
TELECOMMUNICATION INFORMATION:
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                                                                0, Application US/08762308 5925548
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NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:

STREET:

P.O.

Houston

ADDRESSEE:

Arnold, Wh O. Box 4433

White & Durkee

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SEQUENCE CHARACTERISTICS:
LENGTH: 1956 base pairs
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                           110 SerGlnCysValSerSerSerProPheTyrCysGlnProCysLeuAspCysGlyAlaLeu
 492 CAACGCTACCTGAGTGAGACACACTTCCAGTGCGTGGACTGCAGCCCCTGC-----TTC
                                                                                                                                                                                                                                                                                                                                                    210 AAGAGGGATAGCTTGTGTCCCCAAGGAAAGTATGTCCATTCTAAGAACAATTCCATCTGC
                                                                     447 CAAGCTGACAAGGACACGGTGTGTGGCTGTAAG
                                                                                                                                       387 CAGTGTCTCAGTTGCAAGACATGTCGGAAAGAAATGTCCCAGGTGGAGATCTCTCCTTGC
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                                                                                                                                                                        70 GluCysAlaArgCysGlnAlaCysAspGluGlnAlaSerGlnValAlaLeuGluAsnCys
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                       SerAlaValAlaAspThrArgCysGlyCysLysProGlyTrpPheValGluCysGlnVal 109
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US-08-804-166-1
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                                             US-09-993-234-6_COPY_19_204 (1-186) x US-08-804-166-1 (1-1049)
                                                                                                                                                         Pred. No.:
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                                                                                                                                                                                                     ; NAME/KEY:
; LOCATION:
US-08-804-166-1
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INFORMATION FOR SEQ ID NO:
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TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                    FEATURE:
                                                                                                                                                                                                                                                                 MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Browdy, Roger L. REGISTRATION NUMBER: 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 20 February 1996
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                                                                                                                                                                                                                                                                                              STRANDEDNESS:
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HYBRID PROTEINS
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Matches:
Conservative:
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Gaps:
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RESULT 11
US-08-910-991-1
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Patent No. 6194177
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                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                 TELEPHONE: (202) 628-5197
                                                                                                                                                  APPLICATION NUMBER: 08/804,166
FILING DATE: 20 February 1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/011,936
                                                                                                                                                                                                                               CLASSIFICATION: 530 PRIOR APPLICATION DATA:
                                                                                                              FILING DATE: 20 February ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Jameson, Bradford A.
APPLICANT: Chappel, Scott C.
TITLE OF INVENTION: HYBRID PROTEINS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            UMBER OF SEQUENCES:
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                                                                         NAME: YUN, Allen C. REGISTRATION NUMBER:
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                                                           REFERENCE/DOCKET NUMBER:
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                                                                                                                                                                                  Sequence 7, Application Patent No. 6193972
GENERAL INFORMATION:
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                                                                                   APPLICANT: Campbell, Robert K. APPLICANT: Jameson, Bradford A. APPLICANT: Chappel, Scott C. TITLE OF INVENTION: HYBRID PROT NUMBER OF SEQUENCES: 22
                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: BROWDY AND NEIMARK
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MOLECULE TYPE:
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STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                       GCTGCCCCAGGT---TGCCCAGAA 796
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                                                                                                                                                                                                                                                                                                                                                                                                      CysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysProThr 163
                                                                                                                                                                                                                                                                                                                                                                                                                                                           CTCAAT --- GGGACCGTGCAC-----
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Washington D.C.
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible

COUNTRY:

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Best Local Similarity:
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; LOCATION:
US-08-804-166-7
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REFERENCE/DOCKET NUMBER: CAUTELECOMMUNICATION INFORMATION:
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CRASSIFICATION DATA:
PRIOR APPLICATION DATA:
00/011,936
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144 CysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSerCysProThr 163
                                           672
                                                                124 LeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArgAspThrAsp 143
                                                                                                              627 TATTGGAGTGAAAACCTTTTCCAGTGC----
                                                                                                                                                                                567 TCTTCTTGCACAGTGGACCGGGACACCGTGTGTGGCTGCAGGAAGAACCAGTACCGGCAT 626
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                                                                                                                                                                                                                                                                                                67
                                                                                                                                                                                                                                                                                                                                                                  47 CysGlyAsnSerThrCysLeuValCysProGlnAspThrPheLeuAlaTrpGluAsnHis 66
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7 GlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAspPheHisLysLysIleGly
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                                           CTCAAT --
                                                                                                                                                                                                                                                          CTCAGA----CACTGCCTCAGCTGCTCCAAATGCCGAAAGGAAATGGGTCAGGTGGAGATC
                                                                                                                                                                                                                                                                                          HisAsnSerGluCysAlaArgCysGlnAlaCysAspGluGlnAlaSerGlnValAlaLeu
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                                                                                                                                              Cys-----GlnValSerGlnCysValSerSerSerProPheTyrCysGlnProCys 123
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Indels:
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                                                           US-09-993-234-6_COPY_19_204 (1-186) x US-08-910-991-7 (1-1301)
                                                                                                                           Percent Similarity:
Best Local Similarity:
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US-08-910-991-7
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                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (202) 737-35
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 60/0: FILING DATE: 20 February ATTORNEY/AGENT INFORMATION: NAME: YUN, Allen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/804,166
FILING DATE: 20 February 1997
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                                                        MOLECULE TYPE:
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330 CAAGAGGGCAGTGCCGATAGTGTGTGTCCCCAAGGAAAATATATCCACCCTCAAAATAAT 389
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0
                                                                                                                                                                                                                                                       NAME/KEY:
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Matches:
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US-08-050-319B-47
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                                                                                                                                                  CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: ROBDINS, ROBERTA L.
REGISTRATION NUMBER: 33,208
REFERENCE/DOCKET NUMBER: 5150
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 617-8999
                                                                                                                         TELEFAX: (415) 327-32 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                    MOLECULE TYPE: cDNA to mRNA FEATURE:
                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/050,319B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Modified human TNFalpha (Tumor TITLE OF INVENTION: Modified human TNFalpha (Tumor NUMBER OF SEQUENCES: 57
                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: M.Feldmann, P.I
APPLICANT: M.J.C. Turner,
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RESULT 15
US-08-465-982-47
; Sequence 47, Application US/08465982
; Patent No. 5863786
; GENERAL INFORMATION:
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                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                 APPLICANT: M.Feldmann, P.W. Gray,
APPLICANT: M.J.C. Turner, F.M Brennan
TITLE OF INVENTION: Modified human TNFalpha (Tumor
TITLE OF INVENTION: Necrosis Factor alpha) Receptor
                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       139 ArgArgAspThrAspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCys 158
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No.:
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LENGTH: 600 base pairs
TYPE: nucleic acid
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/0:
FILING DATE: 10-May-1993
ATTORNEY/AGENT INFORMATION:
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REFERENCE/DOCKET NUMBER: 5150-0030
TELECOMMUNICATION INFORMATION:
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                          159 ValSerCysProThrSerThrLeuGlySerCysProGlu-----ArgCysAlaAlaVal 176
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SOFTWARE: PatentIn Release #1.0, version #1.25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/465,982
GTCTCCTGT------AGTAACTGTAAGAAAAGCCTGGAGTGCACGAAGTTG
                                                                                                                                         AATTGCAGCCTCTGCCTCAAT---GGGACCGTGCAC-----CTCTCCTGCCAG 477
                                                                                                       ArgArgAspThrAspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCys 158
                                                                                                                                                                                                                  AACCAGTACCGGCATTATTGGAGTGAAAAACCTTTTCCAGTGC-----TTC 432
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Search completed: April 6, 2003, 23:29:58
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-QB-Published_Applications_NA -QEMT=fastap -SUFFIX=rnpb -MINMATCH=0.1
-LOOPCL=0 -LOOPEXT=0 -UNITS-bits -START=1 -END=-1 -MATRIX=blosum62
-TRANS=human40.cdi -LIST=45 -DCCALIGN=200 -THR_SCORE=pct -THR_MX=100
-THR_MIN=0 -ALIGN=15 -MODE=LOCAL -OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0
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-NCPU=6 -ICPU=3 -NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG_
-DEV_TIMEOUT=120 -WARM_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
-FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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Listing first 45 summaries
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Ygapop 10.0 , Ygapext
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/cgn2_6/ptodata/1/pubpna/USOB_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/USO9_NEW_PUB.seq:*
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9 US-10-081-280-9
9 US-10-112-793-9
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Sequence 1, Appli Sequence 1, Appli Sequence 2360, Ap Sequence 26, Appl Sequence 26, Appl Sequence 16, Appl Sequence 16, Appl Sequence 16, Appl	11, 13, 13, 11, 11,	5, e 157 e 157 e 2, e 2, e 2, e 2, e 1, e 1, e 1, e 1,	Sequence 9, Appli Sequence 1, Appli Sequence 1389, Ap Sequence 292, Appli Sequence 5, Appli Sequence 5, Appli

#### ALIGNMENTS

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NESULT 1

US-09-33-966-3

Sequence 3, Application US/09333966

Patent No. US20020009773A1

GENERAL INFORMATION:
APPLICANT: V1, Guo-Liang

APPLICANT: Ni, Jian
APPLICANT: Dixit, Vishya
APPLICANT: Dixit, Vishya
APPLICANT: Dillon, Patrick J.

TITLE OF INVENTION: Death Domain Containing Receptors

NUMBER OF SEQUENCES: 17

CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein & Fox, P.L.L.C.
STREET: 1100 New York Ave., NW, Suite 600

CITY: Washington
STATE: DC
COUNTRY: USA
ZIP: 20005-3934

COMPUTER READABLE FORM:
MEDLUM TYDE: Floppy disk
COMPUTER READABLE FORM:
MEDLUM TYDE: Floppy disk
COMPUTER PATENTIN Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/333,966

FILLING DATE:
CLASSIFICATION:
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 60/028,711
FILING DATE: 17-OCT-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/013,285
FILING DATE: 12-MAR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: 202-371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Steffe, Eric K.
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LOCATION:
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STRANDEDNESS: double
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                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
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                          ASPThrAspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAspGlyCysValSer 160
                                                                                 GlnProCysLeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArg 140
                                                                                                                                        CysProThrSerThrLeuGlySerCysProGluArgCysAlaAlaValCysGlyTrpArg 180
                                                                                                                                                                                              GCCTCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAG
                                                                                                                                                                                                               AlaSerGlnValAlaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLys 100
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 415/952-9881
TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 9:
                                                                                                                                  143
                                                                   203
                  41 AlaProCysThrGluProCysGlyAsnSerThrCysLeuValCysProGlnAspThrPhe 60
                                                                                 21 PheHisLysLysIleGlyLeuPheCysCysArgClyCysProAlaGlyHisTyrLeuLys 40
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                                                                                                                                                1 LeuLeuGlyalaArgAlaGlnGlyGlyThrargSerProArgCysAspCysAlaGlyAsp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: WinPatin (CURRENT APPLICATION DATA:
                                                                                                                            CTGCTGGGGGCCCGGGCCCAGGGCGCACTCGTAGCCCCAGGTGTGACTGTGCCGGTGAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NÜMBER: 35,6
REFERENCE/DOCKET NÜMBER:
TELECOMÜNICATION INFORMATION:
TELEPHONE: 415/225-5416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/829,270 FILING DATE: 31-Mar-1997 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: 3.5 i
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ADDRESSEE: Genentech,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Ashkenazi, Avi
TITLE OF INVENTION: Apo-2
NUMBER OF SEQUENCES: 11
GCCCCTTGCACGGAGCCCTGCGGCAACTCCACCTGCCTTGTGTGTCCCCCAAGACACCTTC
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                                                               TTCCACAAGAAGATTGGTCTGTTTGGTTGCAGAGGCTGCCCAGCGGGGCACTACCTGAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 1634 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 21-Feb-2002 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Marschang, Diane L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/10/081,280
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Matches:
Conservative:
Mismatches:
Indels:
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US-10-112-793-9
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TELEFAX: 650/952-9881
INFORMATION FOR SEO ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 1634 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/112,793
FILING DATE: 28-Mar-2002
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-2 LI AND
NUMBER OF SEQUENCES: 28
    SEQUENCE
                                                                                                                  TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: California
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Best Local Similarity:
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                                                                                                                                                                                                                       Sequence 11, Application US/10112193 Publication No. US20030004313A1
                                                                                                                                                                   GENERAL INFORMATION:
APPLICANY: ASHKenazi, Avi J.
APPLICANY: ASHKENAZi, Avo-3 POLYPEPTIDE
TITLE OF INVENTION: Apo-3 POLYPEPTIDE
NUMBER OF SEQUENCES: 15
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          ZIP: 94080
COMPUTER READABLE FORM:
COMPUTER: 3.5 inch, 1.44 Mb:
MEDIUM TYPE: 3.5 inch, 1.44 Mb:
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT
                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GlnMetPheTrpValGln
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                                                                                                    STREET: 1 DNA Way
CITY: South San F:
STATE: California
                                                                                       COUNTRY:
                                                                             94080
                                                                                        USA
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
                                                 .44 Mb floppy
                                                  disk
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APPLICATION DATA:

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US-09-993-234-6_COPY_19_204 (1-186) x US-10-112-193-11 (1-1634)
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TELEPHONE: 650/225-5416
TELEPAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             143 CTGCTGGGGCCCGGGCCCAGGGCGGCACTCGTAGCCCCAGGTGTGACTGTGCCGGTGAC
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                                                                                               CysProThrSerThrLeuGlySerCysProGluArgCysAlaAlaValCysGlyTrpArg
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FILING DATE: 23-5ep-1996
ATTORNEY/AGENT INFORMATION:
NAME: Marschang, Diane L.
                                                                                                                                                                                                                                            GlnProCysLeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArg
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                                                                              GCCTCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAG
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TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/10/112,193
FILING DATE: 28-Mar-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1052R1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 08/928,069 FILING DATE: 11-Sep-1997
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Matches:
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Indels:
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: 415/225-5416
TELEFAX: 415/952-9881
TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           . No . :
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Wilpatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/993,234
APPLICATION NUMBER: US/09/993,234
CILANG DATE: 19-No. US20020146768A1-2001
CLASSIFICATION: <Unknown>
                                                  TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-5416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 08/828,683
FILING DATE: <UNKNOWN>
FILING GENT INFORMATION:
NAME: MAYSCHANG, Diane L.
REGISTRATION NUMBER: 35,600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Ashkenazi, Avi
TITLE OF INVENTION: Apo-2
NUMBER OF SEQUENCES: 11
               AlaSerGlnValAlaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLys
                                                                                                                                              AlaProCysThrGluProCysGlyAsnSerThrCysLeuValCysProGlnAspThrPhe
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ADDRESSEE: Genentech,
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LENGTH: 1634 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
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CITY: South San Francisco
STATE: California
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Percent Similarity: Best Local Similarity:

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Scores:

No.:

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TOPOLOGY: Linear

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Patent No. US20020009773A1
GENERAL INFORMATION:
                                      INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                     REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 14
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
                                                                                                                                              APPLICATION NUMBER: US 60/013,285
FILING DATE: 12-MAR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Steffe, Eric K.
                                                                                                                                                                                                                         APPLICATION NUMBER: No. US20020009773A1 Yet Assigned FILING DATE: 06-FEB-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/028,711
FILING DATE: 17-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                           FILING DATE: 17-OCT-PRIOR APPLICATION DATA:
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APPLICATION NUMBER:
FILING DATE:
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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    STRANDEDNESS
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1100 New York Ave.,
                               1783 base pairs
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Gentz, Reiner L.
Dillon, Patrick J.
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                                                                                                                                                            Sequence 1389, Application Patent No. US20020165180A1 GENERAL INFORMATION:
                                        APPLICANT: Weaver, Zoe
TITLE OF INVENTION: Process for Identifying
TITLE OF INVENTION: Gene Sets
FILE REFERENCE: 689290-77
CURRENT APPLICATION NUMBER: US/09/954,531
CURRENT FILING DATE: 2002-05-02
PRIOR APPLICATION NUMBER: US/60/233,133
PRIOR FILING DATE: 2000-09-18
             PRIOR APPLICATION NUMBER: US/60/234,009 PRIOR FILING DATE: 2000-09-20
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LOCATION:
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Matches:
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RESULT 8
US-09-964-824A-292
; Sequence 292, Application US/099648
; Patent No. US20020102531A1
; GENERAL INFORMATION:
APPLICANT: HORTIGAN, Stephen
; TITLE OF INVENTION: Cancer Gene De
; TITLE OF INVENTION: Sets
; FILE REFERENCE: 689290-73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 2000-09-20
PRIOR APPLICATION NUMBER: US/60/234,509
PRIOR FILING DATE: 2000-09-22
PRIOR APPLICATION NUMBER: US/60/234,567
PRIOR FILING DATE: 2000-09-22
NUMBER OF SEQ ID NOS: 1392
SOFTWARE: Patentin version 3.0
SEQ ID NO 1389
LENGTH: 1743
TYPE: DNA
ORGANISM: Homo sapiens
US-09-954-531-1389
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Best Local Similarity:
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NUMBER: US/09/964,824A
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                          Determination
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CURRENT FILING DATE: 2001-09-27

PRIOR APPLICATION NUMBER: US/60/236,033

PRIOR FILING DATE: 2000-09-28

PRIOR APPLICATION NUMBER: US/60/236,032

PRIOR APPLICATION NUMBER: US/60/236,032

PRIOR APPLICATION NUMBER: US/60/236,028

PRIOR FILING DATE: 2000-09-28

PRIOR FILING DATE: 2000-09-28

NUMBER OF SEC ID NOS: 583

SOFTWARE: Patentin version 3.0

SEQ ID NO 292

LENGTH: 816

TYPE: DNA

ORGANISM: Homo sapiens

US-09-964-824A-292
                                   RESULT 9
US-10-081-280-5
; Sequence 5, Application US/10081280
; Patent No. US20020165157A1
; GENERAL INFORMATION:
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APPLICANT: Ashkenazi, Avi
TITLE OF INVENTION: Apo-2
NUMBER OF SEQUENCES: 11
                                                                                                                                                    .:::
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Best Local Similarity:
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TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
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No.:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-5416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE DESCRIPTION: SEQ ID
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ADDRESSEE: Genentech,
             GlnProCysLeuAspCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSerArgArg
                                                                                                                                                                                                                                             AlaProCysThrGluProCysGlyAsnSerThrCysLeuValCysProGlnAspThrPhe
                                                                    ProGlyTrpPheValGluCysGlnValSerGlnCysValSerSerSerProPheTyrCys 120
                                                                                                                             AlaSerGlnValAlaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLys
                                                                                                                                                                     TTGGCCTGGGAGAACCACCATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAG
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                                                     CCAGGCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCCTTCTACTGC
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TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
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CITY: South San Francisco
STATE: California
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                                                                     US-09-993-234-6_COPY_19_204 (1-186) x US-10-112-793-5 (1-1438)
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Publication No. US20020192729A1
GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-2 LI
NUMBER OF SEQUENCES: 28
                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 5: SEQUENCE CHARACTERISTICS:
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No.:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/08/828,683A
FILING DATE: 31-Mar-1997
APPLICATION NUMBER: 08/625328
FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
ATTORNEY/ACENT INFORMATION:

NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
MEDIUM TYPE: 3.5 inch, 1.44 Mb COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech) CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 i
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TYPE: Nucleic Acid
STRANDEDNESS: Single
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STREET: 1 DNA Way
CITY: South San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/10/112,793 FILING DATE: 28-Mar-2002
                                                                                                                                                                                                                                                                                  TOPOLOGY: Linear
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Matches:
Conservative:
Mismatches:
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RESULT 11
US-10-112-193-5
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Publicati
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                                                                                                                                            COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 mb floppy dis

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: WinPatin (Genentech)

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/112,193

FILING DATE: 28-Mat-2002

CLASSIFICATION: <Unknown>

PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                              Lication No. US20030004313A1
               APPLICATION NUMBER: 08/928,069
FILING DATE: 11-Sep-1997
APPLICATION NUMBER: 60/026943
FILING DATE: 23-Sep-1996
ATTORNEY/AGENT INFORMATION:
NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REGISTRATION NUMBER: P1052R1
TELEPHONE: 650/225-5416
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TITLE OF INVENTION: Apo-3
NUMBER OF SEQUENCES: 15
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DB:
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Best Local Similarity:
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                                                                         NUMBER OF SEQUENCES:
                                                               CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                 TGCCCCACGTAATTCCTA---
        ADDRESSEE: Genentech, Inc
STREET: 460 Point San Bru
CITY: South San Francisco
STATE: California
                                                                                                                                                                                              949
                                                                                                                                                                                                                        181
COUNTRY: USA
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STRANDEDNESS: Single
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TELEPHONE: 415/225-5416
TELEPAX: 415/952-9811
TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1438 base pairs
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161 CysProThrSerThrLeuGlySerCysProGluArgCysAlaAlaValCysGlyTrpArg 180
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APPLICATION UNMBER: US/09/884,733
FILING DATE: 19-Jun-2001
CLASSIFICATION: CUDKNOWN>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                        81 AlaSerGlnValAlaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLys 100
                                                                                                                                                                                                                                                                                                       61 LeuAlaTrpGluAsnHisHisAsnSerGluCysAlaArgCysGlnAlaCysAspGluGln
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COMPUTER READABLE FORM:
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                                                                                             ProGlyTrpPheValGluCysGlnValSerGlnCysValSerSerSerProPheTyrCys 120
                                                                                                                                                                                                                                                                                                                                                     GCCCCTTGCACGGAGCCCTGCGGCAACTCCACCTGCCTTGTGTGTCCCCAAGACACCTTC
                                               AspThrAspCysG1yThrCysLeuProG1yPheTyrG1uH1sG1yAspG1yCysValSer 160
                                                                                                                                                                                                                           GCCTCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAG
                                                                                                                                                          CCAGGCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGC
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STRANDEDNESS: Single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Marschang, Diane L.
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Matches:
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                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                     COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/993,234
FILING DATE: 19-NO. US20020146768A1-2001
CLASSIFICATION: CUNKNOWN>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007P1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 08/828,683 FILING DATE: <Unknown> ATTORNEY/AGENT INFORMATION:
                                                          PRIOR APPLICATION DATA
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              AlaProCysThrGluProCysGlyAsnSerThrCysLeuValCysProGlnAspThrPhe 60
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TYPE: Nucleic Acid
STRANDEDNESS: Single
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STATE: California
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847-1577 .1577, Application US/09764847 o. US20020132767A1 INFORMATION:		CysGlyTrpArgo	rGluHisGlyAspGlyCysValSerCysProThrSerThrLeuGlySerCysProGluAr 172	CTCCTGGATTGCAGGTTCCCGCAGAGATACTGACTGTGGGACCTGCCTG	jArgAspThrAspCysGlyThrCysLeuProGly	CTACTCCCAGACCCCCTTCTCCCTGCCTGACCCACTCCTGTCCCATGGTGACGCATGCCT 6327	138	CTGCGGGGCCCTGCACCGCCACACACGGCTACTCTGTGA-GTACCCCCCACCCAGGGCTCT 6267	pCysGlyAlaLeuHisArgHisThrArgLeuLeuCysSer 138	62	rCvsGlnProCvsLeuAs 125	dicusiumsuu yseelala valalaaspiilla 1955eltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyyseltyysel	CCCAGGTGGC 60		TCGCCTATTTCTCTCTCCATTATCTTGGGATAATGCCTCTGCCTCTCCATGGGAGCCTTT 6029	79	GGTCGGCCTTAATCTCTGAGCTTCTCTCTTTTTAGGGTAGCCCTGTACCTGTCTGT	79	TGCCCTTGCCCCTTCTCCTTCCTGGTGACCTTCCCATCTCTCCATGTGCCTTGGCCTCT 5909	79	GGGTGTGAGGGTCCTGCACCCACCCTTGCCAGAACCCTCCACCCTGATCCTCCTTCAGGG 5849	79	GCAGGGAGTTCCTAAGGACAGGCCTTTCTGAAGGAAGTGGCTGGC	79	TTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGGGTGAGGGGGCTTCTCAGTGCTTG 5729	sAlaArgCysGlnAlaC		AsnSerThrCysLeuValCysProGlnAspThrPheLeuAlaT		ysGl 48	GAGGGAGGGAGGCAGGTTGGAAAGCAGGTCGGGGGGTTGCTTGGCAGCCCCCTCTGCCTGC	35

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CURRENT FILING DATE: 2001-01-17

Prior application data removed - consult PALM or file wrapper NUMBER OF SEQ ID NOS: 2003

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 1577

LENGTH: 10797

TYPE: DNA

ORGANISM: Homo sapiens
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PC009
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Search completed: April 7, 2003, 03:59:13
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# GenCore version 5.1.4\_p5\_4578 Copyright (c) 1993 - 2003 Compugen Ltd.

Database : EST:*  1: em_esthum:* 2: em_esthum:* 4: em_estov:* 5: em_estov:* 6: em_estov:* 7: em_esti:* 9: qb_esti:* 10: qb_est:* 11: qb_htc:* 12: qb_esti:* 13: qb_esti:* 14: qb_esti:* 15: em_estom:* 15: em_estfun:* 16: em_estfun:* 17: qb_gss:* 18: em_gstfun:* 21: em_gss_nun:* 22: em_gss_pin:* 23: em_gss_pin:* 24: em_gss_other:* 25: em_gss_other:* 27: em_gss_rod:*	Command line parameters:  -MODEL-frame+_p2n.model -DEV=xlh -Q=/cgn2_1/USPT0_spool/US09993234/runat_27032003_115456_15362/app_query.fasta_1.2346 -Q=/cgn7_2_1/USPT0_spool/US09993234/runat_27032003_115456_15362/app_query.fasta_1.2346 -DM=EST -QFMTT-fastap -SUFFIX-rst -MINNATCH+0.1 -LOOPCL-0 -LOOPEXT-0 -UNITS=bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi -LIST-45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_NIN-0 -ALIGN=15 -MODE=LOCAL -OUTEMT=pto -NORM=ext -HEAPSIZE=500 -MAXLEN-200000000 -USER-US09993234_eCGN 1_1_4749_erunat_27032003_115456_15362 -NCPU-6 -ICPU-3 -USER-US09993234_eCGN 1_1_4749_erunat_27032003_115456_15362 -NCPU-6 -ICPU-3 -NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7	Post-processing: Minimum Match 0%  Maximum Match 100%  Listing first 45 summaries	umber of hits sa  DB seq length:  DB seq length:	56 seqs, 809774337	Scoring table: BLOSUM62 Xgapop 10.0 , Xgapext 0.5 Ygapop 10.0 , Ygapext 0.5 Fgapop 6.0 , Fgapoxt 7.0 Delop 6.0 , Delext 7.0	Title: US-09-993-234-6_COPY_19_204 Perfect score: 1101 Sequence: 1 LLGARAQGGTRSPRCDCAGDGSCPERCAAVCGWRQMFWVQ 186	Run on: April 6, 2003, 16:19:54; Search time 1163.45 Seconds (without alignments) 2589.161 Million cell updates/sec	OM protein - nucleic search, using frame_plus_p2n model	Copyright (c) 1993 - 2003 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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## ALIGNMENTS

REFERENCE AUTHORS TITLE		ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION		DEFINITION	LOCUS	BI909448	RESULT 1
<pre>1 (bases 1 to 767) NIH-MGC http://mgc.nci.nih.gov/. National Institutes of Health, Mammalian Gene Collection (MGC)</pre>	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.	Homo sapiens	human.	EST.	BI909448.1 GI:16172666	BI909448	mRNA sequence.	ens cDNA clone In	BI909448 767 bp mRNA linear EST 16-OCT-2001		

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Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (destroyed); RNA source leukocytes from anonymous pool of non-activated adult donors. Library is oligo-dr primed and directionally cloned (EcoRv site is destroyed upon cloning). Average insert size 1.7 kb, insert size range 1.2-3.3 kb. Library is normalized and enriched for full-length clones and was constructed by C. Gruber (Invitrogen). Research Genetics tracking code 027. Note: this is a NIH MGC Library.*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5214568"
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/lab_host="DH10B"
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Location/Qualifiers
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This clone is available royalty-free through
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National Cancer Institute, Cancer Genome Anat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IMAGE Consortium (info@image.llnl.gov) for Insert Length: 1568 Std Error: 0.00
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae.
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                                                                                                                                                                                                                                                                                                                           from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 297480-302087, 682632-687339, 726408-728711, and 729096-731399. Subtraction by E Soares and M. Fatima Bonaldo. " a 218 c 214 g 155 t
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/db_xref="taxon:9606"
/clone="IMAGE:2359204"
/clone_lib="Soares_NFL_T_GBC_S1"
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Contact: Robert Strausberg,
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                                                                                                                                                                                                                                     This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Possible reversed clone: similarity on wrong strand
                                                                                                                                                                                                                                                                                                                              NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy
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Seq primer: -40UP from Gibco
High quality sequence stop: 455.
Location/Qualifiers
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          /note-"Organ: pooled; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Equal amounts of plasmid DNA from three normalized libraries (fetal lung NDHL19W, testis NHT, and B-cell NCI_CGAP_GCB1) were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver
                                                                                                      /clone="IMAGE:2815679"
/clone_lib="Soares_NFL_T_GBC_S1"
/lab_host="DH10B"
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similar to T
Homo saplens
Eukaryota; Metazoa; Chordata; Craniata; Verto
Mammalia; Eutheria; Primates; Catarrhini; Hor
1 (bases 1 to 688)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
                                                               numan .
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                                                                                                            sequence.
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TR:000276 000276 LY
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Craniata; Vertebrata; Catarrhini; Hominidae;

Hominidae;

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                                                                            ProGlyTrpPheValGluCysGlnValSerGlnCysValSerSerSerProPheTyrCys
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Tissue Procurement: Ash Alizadeh, John Byrd, M.D., Mike Greve
M.D., Louis M. Staudt, M.D., Ph.D.
CDNA Library Preparation: M. Bento Soares, Ph.D.
CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing C
Clone distribution: NCI-CGAP clone distribution information
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 2028 Std Error: 0.00
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High quality sequence stop: 445.
Location/Qualifiers
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Unpublished (1997)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone_lib="NCI_CGAP_CLL1"
/tissue_type="B-cell, chronic lymphotic leukemia"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:2109207"
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AUTHORS
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US-09-993-234-6_COPY_19_204 (1-186) x AI140043 (1-562)
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562 bp mRNA linear EST 13-APR-1999
ga87a04.x1 Soares_fetal_heart_NbHH19W Homo sapiens cDNA clone
IMAGE:1693710 3' similar to SW:WSL1_HUMAN Q93038 WSL-1 PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL;
IMAGE Consortium (info@image.llnl.gov) for further j
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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quality sequence stop: 494.
Location/Qualifiers
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a 182
                                                                                                                                                                                                         same fetus as the fetal lung library, Soares fetal lung
                                                                                                                                                                                                                                                                                                                                                                                                                          /sex="unknown"
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/db_xref="taxon:9606"
/clone="IMAGE:1693710"
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1 LeuLeuGlyAlaArgAlaGlnGlyGlyThrArgSerProArgCysAspCysAlaGlyAsp

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Plate: 31 row: H column: 06
High quality sequence stop: 552.
Location/Qualifiers
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52 Eoeun-dong Yuseong-gu, Daejeon 305-333,
                                                                                                                                                                                                                                               Oh, K.J., Cheong, J.E., Kim, Y.S.
                                                                                                                                                                                                                                                                                                                                                               mRNA sequence.
BM783972
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l: +82-42-860-4470
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                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="$6$NU620-31-H06"
/clone_lib="$6$NU620"
/cell_type="Scattering floating"
/cell_line="SNU-620"
                        /tissue_type="Ascites"
                                   /sex-"F"
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E., Sohn,H.Y., Kim,J.M., Park
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circularized with E. coli DNA ligase after digestion of ECORI which site is also included in vector. An RNA strand converted to a DNA strand by Okayama-Berg method. The obtained cDNA vectors were used for transformation of competent cells E. coli ToplOF' by electroporation method. The cDNA libraries constructed by this method are full-length enriched cDNA library."
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BASE

Score: Pred Alignment ORIGIN COUNT

Length: Matches: Conservative:

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585 bp mRNA Soares\_NHCe\_cervix Homo sapiens

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AW517358
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National Cancer Institute, Cancer Genome Anat
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the Not I and Eco RI sites of the modified pT7T3 vector.
Library is normalized; constructed by Bento Soares and
M.Fatima Bonaldo."
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/clone_lib="Soares_NHCe_cervix"
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Fax: +82-42-860-4409
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Plate: 17 row: C column: 06
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/note="Organ: Stomach; Vector: pCNS; Site_1: EcoRI; Site_2: NoLL; The poly (A)+ RNA was dephosphorylated with bacterial alkaline phosphatase (BAP) and then decapped with tabacco acid pyrophosphatase (TAP). The decapped intact mRNA was ligated with DNA-RNA linker including EcoR I site by treatment of T4 RNA ligase and the first strand cDNA was synthesized from oligo dT-selected mRNA by priming with dT-tailed vector. The dT-tailed vector was adjusted to have about 60nt. The cDNA vector was circularized with E. coli DNA ligase after digestion of EcoRI which site is also included in vector. An RNA strand converted to a DNA strand by Okayama-Berg method. The obtained cDNA vectors were used for transformation of competent cells E. coli ToplOF' by electroporation method. The cDNA libraries constructed by this method are full-length enriched cDNA library."
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                Genome Research Center
                                Contact: Kim YS
                                          21C Frontier Korean Unpublished (2002)
                                                                                   Kim, N.S., Hahn, Y., Oh, J.H., Lee, J.Y., Ahn, H.Y.,
Oh, K.J., Cheong, J.E., Sohn, H.Y., Kim, J.M., Park
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                                                                       ProGlyTrpPheValGluCysGlnValSerGlnCysValSerSerSerProPheTyrCys
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Tel: +82-48-680-4470
Fax: +82-42-860-4409
Email: yongsung@mail.kribb
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/clone="56SNU620-27-G03"
/clone_11b="56SNU620"
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Site_2: NotI; The poly (A)+ RNA was dephosphorylated with
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: yongsung@mail.kribb.re.kr
Plate: 32 row: A column: 05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Korea Research Institute of Bioscience & Biotechnology 52 Eceun-dong Yuseong-gu, Daejeon 305-333, South Korea Tel: +82-42-860-4470
Fax: +82-42-860-4409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (20)
Contact: Kim YS
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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m, N.S., Hahn, Y.,
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qe75f02.xl Soares_fetal_lung_NbHL19W Homo sapiens cDNA clone
IMAGE:1744827 3' similar to SW:WSL1_HUMAN Q93038 WSL-1 PROTEIN
                                                                                                                                                                                                                                                                                                                              Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL;
TMAGE Consortium (info@image.llnl.gov) for further
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 508)
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/dev_stage="19 weeks"
/lab_host="DH10B (ampicillin resistant)"
/note="Organ: lung; Vector: pT7T3D (Pharmacia) with
                                                                                                                                                                                                                                                                                                           Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hominidae;
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RESULT 12
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                                                                REFERENCE
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                                                                                                                                                                                                                                                                                         GATACTGACTGTGGGACCTGCCTGCCTGGCTTCTATGAACATGGCGATGGCTGCGTGTCC
71C
                          Kim, N.S., Hahn, Y., Oh, J.H., Lee, J.Y., Ahn, H.Y., Chu, M.Y., Kim, M.I
                                                                           Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                               EST
                                                                                                                                                                             mRNA sequence.
BM741016
                                                                                                                                                                                                          K-EST0013376 S6SNU620 Homo sapiens
                                                                                                             Homo sapiens
                                                                                                                                human.
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                                           (bases 1 to 492)
 Frontier Korean
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           double-stranded cDNA was size selected, ligated to Eco RI adapters (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT7T3 vector (Pharmacia). Library went through one round of normalization to a Cot = 5. Library constructed by Bento Soares and M.Fatima Bonaldo. This library was constructed from the same fetus as the fetal heart library, Soares fetal heart NbHH19W."
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Primates;
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S6SNU620-2-C03 5',
                                             Kim, M.R.,
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Best Local Similarity:
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                                                                                               SerGlnValAlaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLysPro 101
                                                                                                                                                         GCCTGGGAGAACCACCATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGCC
                                                                                                                                                                            AlaTrpGluAsnHisHisAsnSerGluCysAlaArgCysGlnAlaCysAspGluGlnAla
                                                                                                                                                                                                                                      CCTTGCACGGAGCCCTGCGGCAACTCCACCTGCCTTGTGTGTCCCCAAGACACCTTCTTG
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                                                                                                                                                                                                                                                                                                                                      HisLysLysIleGlyLeuPheCysCysArgGlyCysProAlaGlyHisTyrLeuLysAla 41
                                                                          TCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGGCTGTGGCTGTAAGCCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tel: +82-42-860-4470 Fax: +82-42-860-4409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Korea Research Institute of Bioscience & Biotechnology 52 Eoeun-dong Yuseong-gu, Daejeon 305-333, South Korea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   High quality sequence stop: 492
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 circularized with E. coli DNA ligase after digestion of EcoRI which site is also included in vector. An RNA strand converted to a DNA strand by Okayama-Berg method. The obtained cDNA vectors were used for transformation of competent cells E. coli ToplOF' by electroporation method. The cDNA libraries constructed by this method are full-length enriched cDNA library."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note-*Organ: Stomach; Vector: pCNS; Site_1: ECORI; Site_2: NotI; The poly (A)+ RNA was dephosphorylated with bacterial alkaline phosphatase (BAP) and then decapped with tabacco acid pyrophosphatase (TAP). The decapped intact mRNA was ligated with DNA-RNA linker including EcoR I site by treatment of T4 RNA ligase and the first strand cDNA was synthesized from oligo dT-selected mRNA by priming with dT-tailed vector. The dT-tailed vector was adjusted to have about 60nt. The cDNA vector was adjusted to have about 60nt. The cDNA vector was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /cell_type="Scattering
/cell_line="SNU-620"
/lab_host="Top10F'"
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/db_xref="taxon:9606"
/clone="868NU620-2-C03"
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Contact: Kim YS
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                                                                                                                                                                                                                                                                /note-*Organ: Stomach; Vector: pCNS; Site_1: ECORI; Site_2: NOTI; The poly (A)+ RNA was dephosphorylated with bacterial alkaline phosphatase (BAP) and then decapped with tabacco acid pyrophosphatase (TAP). The decapped intact mRNA was ligated with DNA-RNA linker including EcoR I site by treatment of T4 RNA ligase and the first strand cDNA was synthesized from oligo dT-selected mRNA by priming with dT-tailed vector. The dT-tailed vector was adjusted to have about 60nt. The cDNA vector was circularized with E. coli DNA ligase after digestion of EcoRI which site is also included in vector. An RNA strand
                                                                                             obtained cDNA vectors were used for transformation of competent cells E. coli Top10F' by electroporation method The cDNA libraries constructed by this method are full-length enriched cDNA library. After analyzing and sequencing about 2,000 - 3,000 colonies in original cDNA library, the abundant cDNAs were selected and amplified by
                  library, the abundant cDNAs were selected and amplified by PCR reaction using vector region primer including T7 promotor as 5' primer and N(dT)14 as 3' primer. The PCR products were used as template for synthesis of
biotinylated
                                                                                                                                                                                                                                              converted to a DNA strand by Okayama-Berg method. The
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/cell_line="SNU-620"
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/sex="F"
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/db_xref="taxon:9606"
/clone="S6SNU620s1-21-D12"
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ACCESSION VERSION KEYWORDS SOURCE

EST

numan.

BQ068309 BQ068309.1 GI:19897355

sapiens

ORGANISM

Homo sapiens

COMMENT

Unpublished (1999)
Contact: Robert Strausberg, Ph.D.

NIH-MGC http://mgc.nci.nih.gov/. National Institutes of Health, Mammalian

Gene

Collection (MGC)

Euteleostomi;

Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)

REFERENCE

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 1010)

AUTHORS TITLE JOURNAL

DEFINITION

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                                                                                                                   AspThrAspCysGlyThrCysLeuProGlyPheTyrGluHisGlyAsp 156
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                                                                                                     GATACTGACTGTGGGACCTGCCTGCCTGGCTTCTATGAACATGGCGAT
                                                                                                                                                                    CAACCATGCCTAGACTGCGGGGGCCCTGCACCGCCACACGCCTACTCTGTTCCCGCAGA
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1 156 c 147 g 1
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AW002222 478 bp mRNA linear ES: wt85c09.xl Soares_thymus_NHFTh Homo sapiens cDNA clone IMAGE:2514256 3' similar to TR:014866 014866 SOLUBLE DI
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Plate: LLAM12833 row: n column:
High quality sequence stop: 689.
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/lab_host="DH10B"
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101°ProGlyTrpPheValGluCysGlnValSerGlnCysValSerSerSerProPheTyrCys
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                                                         AlaSerGlnValAlaLeuGluAsnCysSerAlaValAlaAspThrArgCysGlyCysLys 100
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                                                                                                                                                                     AlaProCysThrGluProCysGlyAsnSerThrCysLeuValCysProGlnAspThrPhe
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                                                                                                   TTGGCCTGGGAGAACCACCATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAG
                                            GCCTCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAG
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High quality sequence stop: 410.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL;
IMAGE Consortium (info@image.llnl.gov) for further:
Insert Length: 1648 Std Error: 0.00
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National Cancer Institute, Cancer Genome Anat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AW002222
AW002222.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert Strausberg, Ph.D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tumor Gene Index
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /lab_host-"DH10B (phage-resistant)"
/note-"Organ: thymus, pooled; Vector: pT7T3D-Pac
(Pharmacia) with a modified polylinker; Site_1: t
Site_2: Eco RI; 1st strand cDNA was primed with a
coligo(dT) primer [5'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1. .478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone_lib="Soares_thymus_NHFTh"
/dev_stage="fetal"
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Command line parameters:

-MODEL-frame+_p2n.model -DEV-xlh
-Q-cgn2_flVSPT0_spool/US09993234/runat_27032003_115455_15349/app_query.fasta_1.2346
-DB-GenEmb1 -QFMT-fastap -SUFFIX=rge -MINMATCH-0.1 -LOOPCL-0 -LOOPEXY=0
-UNITS-blits -STARY-1 -END-1 -MATRIX-blosum62 -TRANS-human40.cdi -LIST-45
-DOCALIGN-200 -THR_SCORE-pct -THR_MAX-100 -THR_MIN-0 -ALIGN-15 -MODE-LOCAL
-OUTFMT-pto -NORM-ext -HEAPSIZE-500 -MINLEN-0 -MAXIEN-200000000
-USER-US0999234 @CGN 1_1_8534_@runat_2703203_115455_15349 -NCPU-6 -ICPU-3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NG_SCORES-0 -WAIT -LONGLOG -DEV_TIMEOUT-120
-WARN_TIMEOUT-30 -THREADS-1 -XGAPOP-10 -XGAPEXT-7
-YGAPOP-10 -YGAPEXT-0.5 -DELOP-6 -DELEXT-7
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Maximum Match 100%
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                             ALIGNMENTS
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                                                                                                                                       AF329969 Mus muscu
AF134858 Mus muscu
G37503 SHGC-57889
M59377 Murine tumo
U19994 Sus scrofa
M60468 Mouse tumor
X59238 Murine mRNA
BC004599 Mus muscu
L26349 Mus musculu
                                                                                                                                                                                                                                                                                       AL158217 Human DNA
AF026070 Homo sapi
AX331947 Sequence
Y09392 H.saplens m
AF026071 Homo sapi
BC017526 Mus muscu
AC118359 Rattus no
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U94501 Human 1ymph
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U94503 Human 1ymph
U84503 Human death
U75380 Human apopt
AK094463 Homo sap1
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AB051851 Homo sapi
AB051850 Homo sapi
AL158217 Human DNA
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AB051103 Felis cat
AF329976 Rattus no
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AR119657 Sequence
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U94510 Human lymph
U94504 Human lymph
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      586
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AlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGluMet 40
                                                                GCAGAGATCGAAGCCGTGGAGGTGGAGATCGGCCGCTTCCGAGACCAGCAGTACGAGATG
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Submitted (19-MAR-1997) Molecular Immunology
Submitted Medicine, John Radcliffe Hospital,
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LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
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Accession Number U94501, probable skipping
exons 3, 4, 5, 6 and 7"
236 c 248 g 109 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /product="lymphocyte associated receptor of death 7"
/protein_id="AAC51313.1"
/db_xref="G1:2071963"
/t-anslation="MEQREPROCAVAAALLLVLLGARAQGGTRSPRCDCAGDEHKKIG
LFCCRGCPADEAGMEALTPPEATHLSPIDSAHTLLAPPDSSEKICTVQLVGNSWTPGY
PETQEALCPQVTWSWDQLPSRALGPAAAPTLSPESPAGSPAMMLQPGPQLYDVMDAVP
ARRWEETVRTLGLREBEIEAVEVEIGRFRDQQYEMLKRWRQQQPAGLGAVYAALERMG
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/note="similar to Fas and TNF-R1; lacks transmembrane
/note="similar to Fas and TNF-R1; lacks transmembrane
domain; possibly soluble; LARD-6; LARD-7; NGFR family
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
/db_xref="taxon:9606"
/cell_line="HeLa"
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1 MetAspAlaValProAlaArgArgTrpLysGluPheValArgThrLeuGlyLeuArgGlu
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Direct Submission
Submitted (19-MAR-1997) Molecular Immunology
Submitted (19-MAR-1997) Molecular Immunology
Submitted (19-MAR-1997) Molecular Immunology
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LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
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                                                                                                                                                                                                                                                                                                                              exon 3"
328. .329
                                                                                                                                                                                                                                    /note-"deletion compared to LARD-1a, deposited in GenBank Accession Number U94501, probably due to skipping of putative exons 5, 6 and 7, leading to premature transcriptional termination"

281 c 285 g 138 t
                                                                                                                                                                                                                                                                                                                                                                                                                         /product="lymphocyte associated receptor of death 6"
/protein_id="AAC51312.1"
/db_xref="G1:2071961"
/translation="MEORPRGCAAVAAALLLVLLGARAQGGTRSPRCDCAGDFHKKIG
LFCCRCCPAASQVALENCSAVADTRCGCKPGWFVECQVSQCVSSSPFYCQPCLDCGAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /function="mediates apoptosis"
/note="similar to Fas and TNF-R1; truncated before transmembrane domain; possibly soluble; LARD-6"
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Mismatches:
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Submitted (19-MAR-1997) Molecular Immunology
Submitted Medicine, John Radcliffe Hospital,
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 952)
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                                                                                                     /note-"deletion compared to LARD-1a, deposited in Accession Number U94501, probably due to skipping putative exons 6 and 7, leading to premature transcriptional termination"

317 c 319 g 162 t
                                                                                                                                                                                    exon 3"
407. .408
                                                                                                                                                                                                                                                           /translation="MEQRPRGCAAVAAALLLYLLGARAQGGTRSPRCDCAGDFHKKIG
LFCCRGCPAASQVALENCSAVADTRCGCKPGWFVECQVSQCVSSSPFYCQPCLDCGAL
HRHTRLLCSRRDTDCGTCLPGFYEHGDGCVSCPT"
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soluble; truncated before the
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Accession Number U94501,
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/protein_id="AAC51311.1"
/db_xref="GI:2071959"
                                                                                                                                                                                                                                                                                                                                                                                      /function="mediates apoptosis"
/note="LARD-5; similar to Fas
                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Homo sapiens"
/db_xref="taxon:9606"
/cell_line="HeLa"
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                                                                                                                                                                                                                                                                                                                                                                                                                                     /cell_type="lymphocyte"
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A, alternatively
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U94505
U94505.1
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Screaton,G.R., Xu,X.N., Olsen,A.L., Cowper,A.E., Tan,R.,
McMichael,A.J. and Bell,J.I.
LARD: a new lymphoid-specific death domain containing receptor
regulated by alternative pre-mRNA splicing
Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
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lymphocyte associated receptor of death
                                exons 6 and 7 leading to premature transcriptional termination"
                                                                                                                                                                                                                                                       /function="mediates apoptosis"
/note="LARD-4; similar to Fas
soluble; truncated before the
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                                                                 /note="deletion compared to LARD la,
Accession Number U94501, probable ski
                                                                                                                    /translation="MEORPRGCAAVAAALLLVLLGARAQGGTRSPRCDCAGDFHKKIG
LFCCRGCPAGHYLKAPCTEPCGNSTCLVCPQDTFLAWENHHUSECARCQACDEQASQV
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                                                                                                                                                                                     /product="lymphocyte associated receptor
/protein_id="AAC51310.1"
/db_xref="GI:2071957"
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1. .1087
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/db_xref="taxon:9606"
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spliced, complete cds
U94509
U94509.1 GI:2071964
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Direct Submission
Submitted (19-MAR-1997) Molecular Immunology Group, Institute
Molecular Medicine, John Radcliffe Hospital, Headington, Oxfor
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LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing
Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 1119)
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               PPDSSEKICTYQLYGNSWTPGYPETQEALCPQVTWSWDQLPSRALGPAAAPTLSPESPAGSFAMMLQPGPQLYDYMDAVPARRWKEFVRTLGLREAEIEAVEVEIGRFRDQQYEML
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/product="lymphocyte associated
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                                                                                                                                                        TNF-R1"
                                                                                                                                                                /function="mediates apoptosis"
/note="LARD-8; NGFR family member;
                                                                                                                                                                                                                        /cell
                                                                                                                                                                                              /cell_type="lymphocyte"
1. .1119
                                                                                                                                                                                                                                    /organism="Homo sapiens"
/db_xref="taxon:9606"
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US-09-993-234-6_COPY_339_409 (1-71) x HSU94509 (1-1119)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTCAAGCGCTGGCGCCAGCAGCCAGCCCGCGGGCCTTCGGAGCCGTTTACGCGGCCCTGGAG
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                                                                                                                                                                                                                                                                                                                        Submitted (19-MAR-1997) Molecular Immunology Group, Institute of Molecular Medicine, John Radcliffe Hospital, Headington, Oxford OX3
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Screaton,G.R., Xu,X.N., Olsen,A.L., Cowper,A.E., Tan,R., McMichael,A.J. and Bell,J.I.
LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997) 97272273
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human lymphocyte associated receptor spliced, complete cds. U94510
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/product="lymphocyte associated receptor of death
/protein_id="AAC51315.1"
/db_xref="GI:2071967"
                                                                                       TNF-R1"
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/note="deletion compared to LARD 1a, deposited in GenBank Accession Number U94501, probable skipping of putative exon 3"
1 374 c 371 g 199 t
                                                                                                         /function="mediates apoptosis"
/note="LARD-9; NGFR family mem
                                                                                                                                                                             /cell
                                                                                                                                                                                               /db_xref="taxon:9606"
/cell_line="HeLa"
                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                 /codon_start=1
                                                                                                                                                                                                                                            'organism="Homo sapiens'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GI:2071966
                                                                                                                                                    .eii_type="lymphocyte"
.1143
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Indels:
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or of death 9 mRNA, alternatively
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                                                                                                                                                                                                              Mammalia; Lucinia (198)

1 (bases 1 to 1198)

Screaton,G.R., Xu,X.N., Olsen,A.L., Cowper,A.E., Tan,R.,
McMichael,A.J. and Bell,J.I.

LARD: a new lymphoid-specific death domain containing receptor 'regulated by alternative pre-mRNA splicing
regulated by alternative pre-mRNA splicing
Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                     Human lymphocyte associated receptor of death 3 mRNA, alternatively spliced, complete cds.
                                                                                                          Direct Submission

Submitted (19-MAR-1997) Molecular Immunology Group, Institute of Molecular Medicine, John Radcliffe Hospital, Headington, Oxford OX3
                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
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DSAHTLLAPPDSSEKICTVQLVGNSWTPGYPETQEALCPQVTWSWDQLPSRALGPAAA
PTLSPESPAGSPAMMLQPGPQLYDVMDAVPARRWKEFVRTLGLREAEIEAVEVEIGRF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /translation="meqrprgcaavaaalllyllgaraqggtrsprcdcagdfhkkig
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ALENCSAVADTRCGCKPGWFVECQVSQCVSSSPFYCQPCLDCGALHRHTRLLCSRRDT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note="deletion compared to LARD 1a, deposited in GenBank Accession Number 194501, probable skipping of putative exon 7 encoding the transmembrane domain" 378 c 378 g 199 t
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/cell_type="lymphocyte"
1. .657
                                                                                             Location/Qualifiers
                                                /db_xref="taxon:9606"
                                                              organism="Homo sapiens"
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RESULT 7 HSU94504 LOCUS

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Tittle,T.V. and Wegmann,K.W.
Tr3-specific binding agents and methods fo Patent: WO 0135955 A 125-MAY-2001;
Tittle, Thomas V. (US); Wegmann, Keith W. Location/Qualifiers
20 1. 1250
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/note="deletion compared to LARD la, deposited in GenBank Accession Number U94501, probable skipping of putative exon 6 leading to premature transcriptional termination" a 405 c 384 g 216 t
                                                                                                                                                                                                                                                               /organism="Homo sapiens"
/db_xref="taxon:9606"
420 c 406 g 22:
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DCGTCLPGFYEHGDGCVSCPTVLGPGAPGWPCGPPPAWGHPDLHIPPLLASQAPGYCR
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soluble; truncated before the transmembrane
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/note="LARD-3; similar to Fas
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Signal transduction by DR3, a death domain-containing receptor related to TNFR-1 and CD95
Science 274 (5289), 990-992 (1996)
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VGNSWTPGYVEDTADEAGMEALTPPPATHLSPLDSAHTLLAPPDSSEKICTVQL
VGNSWTPGYVETCGALCPQVTWSWDQLDSRALGFAAAFTLSPESPAGSFAMMLQPGPQ
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Submitted (19-MAR-1997) Molecular Immunology Group, Institute of
Molecular Medicine, John Radcliffe Hospital, Headington, Oxford OX3
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LYDVMDAVPARRWKEFVRTLGLREAEIEAVEVEIGRFRDQQYEMLKRWRQQQPAGLGA
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ALENCSAVADTRCGCKPGMFVECQVSGCVSSSFFYCQPCLDCGALHRHTRLLCSRRDT
DCGTCLPGFYEHGDGCVSCPFSTLGSSCPERCAAVCGMRQMFWYQVLLAGLVVPLLLG
TLTYTYRHCMPHKPLVTADEAGMEALTPPPATHLSPLDSAHTLLAPPDSSEKICTVQL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /function="mediates apoptosis"
/note="LARD-la; membrane prote
TNF-R1; contains a death domai
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/protein_id="AAC51306.1"
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/translation="MEQRPRGCAAVAAALLLVLLGARAQGGTRSPRCDCAGDFHKKIG
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                                                                      GCAGAGATCGAAGCCGTGGAGGTGGAGATCGGCCGCTTCCGAGACCAGCAGTACGAGATG
                                                                                       AlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGluMet 40
CTCAAGCGCTGGCGCAGCAGCAGCCGGGGCCTCGGAGCCGTTTACGCGGCCCTGGAG
                   LeuLysArgTrpArgGlnGlnProAlaGlyLeuGlyAlaValTyrAlaAlaLeuGlu 60
                                                                                                                                             ATGGACGCGCTCCCAGCGCGCGCGCTGGAAGGAGTTCGTGCGCACGCTGGGGCTGCGCGAG
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Accession Number U94501, probably repr
3' splice site"
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LVGNSWTPGYPETQEALCPQYTWSWDQLPSRALGPAAAPTLSPESPAGSPAMMLQPGP
QLYDVMDAVPARRWKEFYRTLGLREAEIEAVEYEIGRFRDQQYEMLKRWRQQQPAGLG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /translation="MEQRPRGCAAVAAALLLYLLGARAQGGTRSPRCDCAGDEHKKIG LFCCRGCPAGHYLKAPCTEPCGNSTCLVCPQDTFLAWENHHNSECARCQACDEQASQV ALENCSAVADTRCGCKPGWFVECQVSQCVSSSPFYCQPCLDCGALHRHTRLLCSRRDT DCGTCLPGFYEHGDGCVSCPTSTLGSCPERCAAVCGWRQMFWVQVLLAGLVVPLLLGA
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TNF-R1; contains a death domain"
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/protein_id="AAC51307.1"
/db_xref="GI:2071951"
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21 AlaGluILeGluAlaValGluValGluILeGlyArgPheArgAspGlnGlnTyrGluMet 40
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Submitted (19-MAR-1997) Molecular Immunology Group, Institute of
Molecular Medicine, John Radcliffe Hospital, Headington, Oxford
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     l (bases 1 to 135)
Screatron,G.R., Xu,X.N., Olsen,A.L., Cowper,A.E., Tan,R.,
McMichael,A.J. and Bell,J.I.
LARD: a new lymphoid-specific death domain containing receptor
regulated by alternative pre-mRNA splicing
Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
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U94503
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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                                                                                                                                                                                                                                                                                                                          /note="insertion compared to LARD la, deposited in GenBank Accession Number U94501, probable retained intron leading to premature transcriptional termination" 441 c 451 g 240 t
                                                                                                                                                                                                                                                                                                                                                                                                                      ALENCSAVADTRCGCKPGWFVECQVSQCVSSSPFYCQPCLDCGALHRHTRLLCSRRDT
DCGTCLPGFYEHGDGCVSCPTSTLGSCPERCAAVCGWRQSRWCAGNASGRTGMDRGEA
GEEGGNHPTPTSCFQCSGSRCSWLALWSPSCLGPP"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /product="lymphocyte associated receptor of death 2"
/protein_id="AAC51308.1"
/db_xref="G01:2071953"
/translation="MEORPRGCAAVAAALLLVLLGARAQGGTRSDRCDCAGDEHKKIG
/translation="MEORPRGCAAVAAALLLVLLGARAQGGTRSDRCDCAGDEHKKIG
/translation="MEORPRGCAAVAAALLLVLLGARAQGGTRSDRCDCAGDEQASQV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note="LARD-2; similar to Fas and TNF-R1; possibly soluble; truncated before the transmembrane domain
                                                                                                                                                                                                                                                                                                                                                                                                           599. .700
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  function="mediates apoptosis"
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/db_xref="taxon:9606"
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                                                                                    LeuLysArgTrpArgGlnGlnGlnProAlaGlyLeuGlyAlaValTyrAlaAlaLeuGlu 60
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                                                                    ATGGACGCGGTCCCAGCGCGCGCGCTGGAAGGAGTTCGTGCGCACGCTGGGGGCTGCGCGAG
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( bases 1 to 1528)

Chaudhary, P.M. and Hood, L.E.

Direct Submission

Submitted (03-JAN-1997) Molecular Biotechnology, University of Submitted (03-JAN-1997) Molecular Biotechnology, WA 98195, ...
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                                                                                                                                                                                                                                                                                                                                                              /translation="GCAAVAAALLLVLLGARAQGGTRSPRCDCAGDFHKKIGLFCCRG
CPAGHYLKAPCTEPGCNSTCLVCPQDTFLAWENHHNSECARCQACDEQASQVALENCS
AVADTRCGCKFGWFVECQVSGCVCSSSPYCQPCLDCGALHRHTLLGSRDTDCGTCL
LGFYEHGDGCVSCPTSTLGSCPERCAAVCGWRQMFWVQVLLAGLVVPLLLGATLTYTY
RHCWPHKPLVTADEAGWEALTPPPATHLSPLDSAHTLLAPDDSSEKICTVQLVGNSWT
PGYPETQEALCPQVTWSWDQLPSRALGPAAAPTLSPESPAGSPAMMLQPGPQLYDVMD
AVPARRWKEFVETLGLEAEE EAVEVEIGRFRDQQYEMLKRWRQQQPAGLGAVYAALE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /function="mediates apoptosis when cross-linked"
/functe="A TNFR1-related death-domain containing receptor;
/note="A TNFR1-related death-domain containing receptor;
DR3, WSL-LR, Apo-3, TRAMP; transmembrane form; similar to
human EST clone 298913, GenBank Accession Number N71143"
                                                                                                                                                                                                                                                                                                                                   RMGLDGCVEDLRSRLQRGP"
496 c 470 g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /product="death domain receptor 3"
/protein_id="AAB41432.1"
/db_xref="GI:1800293"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /gene="DDR3"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /map="1p36"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /chromosome="1"
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Oy 41 LeuLysargTrpArgGlnGlnProAlaGlyLeuGlyAlaValTyrAlaAlaLeuGlu 60
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Search completed: April 6, 2003, 21:08:13 Job time : 829.126 secs

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-Q-/cgn2_1/USPTO_spool/US09993234/runat_27032003_115454_15312/app_query.fasta_1.2346
-DB-N_Geneseq_101002 -QFMT=fastap -SUFFIX=rng -MINKATCH=0.1 -LOOPCL=0
-LOOPEXT=0 -UNITS-bits -START=1 -END=-1 -MATRXX=blosum62 -TRANS=human40.cdi\
-LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MIN=0 -THR_MIN=0 -ALIGN=15
-MODE-LOCAL -OUTFMT=pto -NORM-ext -HEASIZE=500. —NINLEN=0 -MAXLEN=200000000
-USER=US09993234 @CGN_1_1_660_@runat_27032003_115454_15312 -NCPU=6 -ICPU=3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPEXT=7
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Pred. No. is th score greater t and is derived is the number of results predicted by chance to have a ster than or equal to the score of the result being printed, rived by analysis of the total score distribution.

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2161 2170 2175 2176 2176 2254 441 1062 1062 1236	511 2130 2173 1368 1368 2062 2088 2088 2111 2111 2111 2111 2111 211	1250 1254 1254 1254 1254 1634 1634 1662 1783 1783 1783 1783 1783 1797 1797 1797 1797 1793 1793 1793 179	Length
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ABK13194 AAQ50870 AAQ90513 AAQ12215 AAQ8104 AAA851104 AAA188626 AAY63095 AAC84744 AAX23721	ABQ47533 ABK635694 AAQ06284 AAQ06284 AAQ49932 AAA95105 AAQ224440 AAQ210883 AAQ210883 AAQ210955 AAZ09170 AAH48859 AAX09170 AAH48859 ABK64039 ABK64039 ABK64039 ABK64039 ABK64039 ABK64039 ABK64039 ABK64039	AAF83770 AAT89427 AAX00925 AAC68777 AAT91180 AAH27186 AAC91477 AAC91477 AAC9147186 AAC91477 AAC9147531 ABC47531 ABC47531	SUMMARIES
Human tumour necrosi p55 Tumour necrosi p55 TNF-R gene. H Type I TNF recepto Partial human TNFR Human polynucleoti Human TR6 partial Partial nucleotide Human killer adria	Oligonucleotide for Rat sequence differ Rat Tumour Necrosi Lambda-derived TNF Human TNFR1 coding TNF-alpha binding Encodes TNF-alpha 30kD TNF inhibitor Human 30kD TNF inhibitor Human TNFBP-associ Human TNFBP-associ Human CDNA differe Gene #2360 used to Human Tumour necro Human Tumour necro	Nucleotide sequence Death domain conta Death domain conta Human death domain Human apoptosis pr Human genomic DNA. Human pro779 cDNA. cDNA encoding human Death domain conta Death domain conta Death domain conta Human Dray gene ass Genomic sequence # Breast cancer rela Mouse apoptosis in Oligonucleotide fo Oligonucleotide fo	cription

## RESULT 1 AAF83770

ALIGNMENTS

AAF83770 standard; DNA; 1250 BP.

## AAF83770;

06-AUG-2001 (first entry)

Nucleotide sequence of human TR3 gene.

TR3; cell proliferation; leukemia; immunosuppressive; cytostatic; dermatological; antiarthritic; antidiabetic; neuroprotective; car antithyroid; antiinflammatory; antiallergic; T-cell-inhibitor; ds cardiant;

## Homo sapiens

Location/Qualifiers
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                                                                                                 US-09-993-234-6_COPY_339_409 (1-71)
                                                                                                                                                                                                                                        Query Match:
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Best Local Similarity:
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                                                                                                                                                                                                                                                                        TR3-specific binding agent (I) that binds to TR3 and inhibits the proliferation of cells expressing TR3. (I) identified by the methods are useful for treating a subject suspected of having a disease associated with a proliferation of cells expressing TR3 especially leukemias or lymphomas or a T-cell mediated disease especially autoimmune diseases such as myasthenia gravis, systemic lupus erythamatosus, rheumatoid arthritis, diabetes, multiple sclerosis, sarcoidosis, myocarditis, thyroiditis and tumours. (I) is also useful for treating a subject suspected of having graft-versus-host disease, rejection of a transplanted organ such as heart, liver, lung, kidney, pancreas, bowel, skin or an appendage, or inflammatory diseases, allergies and contact dermatitis. The present sequence represents the nucleotide sequence of
                                             1011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mat_peptide
                                                                                                                                                                                                                                        Sequence 1250
                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a composition comprising a biologically active
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 72; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-343711/36.
P-PSDB; AAB84941.
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AlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGluMet 40
                                      TR3 gene.
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/*tag= c
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                                  This cDNA clone codes for human death domain containing receptor 20 DR3 (see AAW31517), a novel member of the tumour necrosis factor receptor family. It was isolated from a HUVEC cDNA library. Related death domain containing receptor DR3-VI cDNA (see AAT89426) was isolated from a human testis tumour cDNA library. The genes have also been identified in cDNA libraries of foetal liver, foetal brain, tonsil and leukocyte. Nucleic acids encoding full-clength or mature DR3, or the extracellular, transmembrane, intracellular or especially the death domain of DR3, can be used to produce recombinant polypeptides in transformed host cells. These polypeptides can be used to treat diseases and disorders associated with the inhibition of apoptosis. Antagonistscan be used to treat diseases and disorders associated with increased apoptosis and for treating inflammatory alsoace and disorders associated with the inhibition of apoptosis.
                                                                                                                                                                                                                                          Claim 6; Page 75-77; 108pp; English.
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                                                                                                                                                                                                                                                                      for activation of apop
to treat inflammatory
                                                                                                                                                                                                                                                                                  Death domain containing receptor polypeptide(s) DR3 and DR3-V1 - for activation of apoptosis and NF-kappaB, antagonists can be used
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(UNMI) UNIV MICHIGAN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                inflammation;
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DB; AAW31517.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            domain
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                        and disorders associated with inflammatory diseases and
                                                                                                                                                                                                                                                                                                                                                                           Dixit VM,
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Pred. No.: Alignment Scores:

2.24e-40 368.00

Length: Matches:

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Sequence

1254

BP;

201 A; 420 C;

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        The invention provides nucleotide sequences encoding death domain containing receptor polypeptides DR3-V1 and DR3. The DR3-V1 cDNA clone is contained in ATCC deposition No. 97456 and the DR3 cDNA clone is contained in ATCC deposition No. 9757. Recombinant vectors comprising the nucleic acid sequences and optionally the leader sequences are
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12-MAR-1996;
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                                                                                                         death domain containing
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UNIV MICHIGAN.
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                                                                                                                                                                                                                                                                                                                                                                                                                          receptor; DR3-V1;
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96US-0013285.
96US-0028711.
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production
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                                                      Yu G,
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28-MAY-1999;
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                                                                                                                                                                                                                   Human; death
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                                                                                                                                                                                                                                                                                                            ArgMetGlyLeuAspGlyCysValGluAspLeu
                                                                                                                                                                                                                                                                                                                                CTCAAGCGCTGGCGCCAGCAGCAGCCCGCGGGCCTTCGGAGCCGTTTACGCGGCCCTGGAG
                                 2000-687263/67.
DB; AAB36265.
                                                     Ni J,
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                                                                    DIXIT
                                                                          GENTZ R L.
DILLON P J.
                                                                                                              HUMAN
                                                                                                      UNIV MICHIGAN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1254 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  represents a cDNA encoding polypeptide (DR3).
                                                                                                                                                                                                                                                                         standard;
                                                                                                                                                                                                          domain containing receptor; DR3; cancer;
isorder; inflammation; cardiovascular disorder; infection;
                                                                                                                                              2000WO-US10741
                                                                                                                                                                                                                                             (first entry)
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                                                                                                             GENOME SCI INC
                                                      Gentz
                                                                                                                          99US-0130488.
99US-0136741.
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368.00
100.00%
y: 100.00
100.0
                                                                                                                                                                                                     disease; angiogenesis; ss.
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Matches:
Conservative:
Mismatches:
Indels:
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Treating graft-versus-host disease, cancer, immunodeficiency or autoimmune disease comprising administering an antibody to Death Containing Receptor proteins and a second therapeutic agent -

Domain

Example

2в;

Fig

<u>ب</u>

273pp;

English.

The present invention provides the protein and coding sequences for two death domain containing receptors, designated DR3 and DR3-V1. These receptors are involved in apoptosis, and the sequences given can be used in the treatment of cancers, infections, cardiovascular disorders such as arrhythmias, ischaemia, aneurysms, arterial occlusive diseases, embolisms and congenital heart defects, neurodegenerative diseases including Alzhelmer's and Parkinson's disease, autoimmune disease such as multiple sclerosis, arthiltis, diabetes, Graves' disease, asthma and psoriasis,

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RESULT 5
AAT91180
ID AAT9
XX AAT9
AC AAT9
AC AAT9
AX Huma
XX Apo-
XX Homo
XX Homo
CS Homo
CFT CDS
FT CDS
FT Sig_
FT mat_
FT mat_
YX XX
XX
XX
XX
PN W097
XX W097
XX 31-M
XX 23-S
PR 23-S
PR 23-S
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                                                   31-MAR-1997;
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                    23-SEP-1996;
01-APR-1996;
                                                                        09-OCT-1997.
                                                                                            W09737020-A1
                                                                                                                        mat_peptide
 (GETH ) GENENTECH INC
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                                                                                                                                                                                                                               Apo-3; apoptosis;
                                                                                                                                                                                                                                                   Human apoptosis protein Apo-3 cDNA clone FH20.57
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                   96US-0710802
96US-0625328
                                                   97WO-US05230
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89..160
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/note= "determined
161..1339
                                                                                                                                                                          Location/Qualifiers
89..1342
                                                                                                               ′*tag=
                                                                                                                                                                                                                              human; therapy; drug screening;
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RESULT 6
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Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Alignment Scores: Pred. No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cDNA clone FH20.57 (ATCC 55820) codes for a novel polypeptide (see w26709), designated Apo-3, that stimulates or induces apoptotic activity in mammalian cells. It was isolated from a human foetal heart cDNA library by screening with probes (see T91183-84) based on an EST sequence (GenBank locus W71994) that showed homology to the intracellular domain of human TNFR1 and CD95. Amino acid residues 1-181 of Apo-3 are identical to another novel apoptosis polypeptide, Apo-2LI (see W26708). Nucleic acids encoding Apo-3 can be used diagnostically for tissue-specific typing and to produce recombinant Apo-3 polypeptides, especific typing and to produce recombinant Apo-3 polypeptides, especificly the extracellular domain (amino acids 1-198) or death domain (amino acids 38-417). Apo-3 can be used to induce apoptosis or NF-kappa-N- or JNK-mediated gene expression for therapeutic purposes. Non-human transgenic animals containing cells that express Apo-3 nucleic acid, and knockout animals containing cells that cells that have an altered Apo-3 gene, can be used in drug screening and development.
                 CDS
                                                                                                                                                                                                                                                                       1283
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                                                                                                                 Human genomic DNA encoding a rheumatoid arthritis associated
                                                                                                                                                                                                                                                                                                                                                                                            1163
                                                                                        Rheumatoid arthritis;
                                                                                                                                                15-AUG-2001
                                                                                                                                                                               AAH27782;
                                                                                                                                                                                                            AAH27782 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1634 BP; 300 A; 528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        screening and development.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 45-46; 70pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polypeptide(s) Apo-3 and Apo-2LI - useful for regulating apoptosis in mammalian cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1997-503105/46.
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                                                          sapiens
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                                                                                                                                              (first entry)
               Location/Qualifiers 89..1342
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Indels:
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41

LeuLysArgTrpArgGlnGlnGlnProAlaGlyLeuGlyAlaValTyrAlaAlaLeuGlu

1134 40 1074

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US-09-993-234-6\_COPY\_339\_409 (1-71) x

AAC68777 (1-1254)

Percent Similarity:
Best Local Similarity:
Query Match:

2.24e-40 368.00 100.00% 100.00% 100.00%

Length:
Matches:
Conservative: Mismatches: Indels:

1254 71 0

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Alignment Scores: Pred. No.:

No . :

Sequence 1254 BP;

to promote angiogenesis and wound healing.

201 A; 420 C; 407

G; 226 T;

0 other;

DB:

В

SQXCCCCCXXXXIII

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RESULT 7
AAL47186
ID AAL4
XX
AC AAL4
XX
DT 22-A
XX
DT 22-B
UMB
XX
KW Huma
XX
KW Huma
XX
OS Homc
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Query
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                                                                                                                                                                                                                                                                                                                                          Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                  This invention relates to a method of diagnosing chronic rheumatoid arthritis by digesting human genomic DNA with EcoRI and hybridising with a probe containing a fragment of the present sequence which represents DNA encoding a transmembrane protein. The method is used the diagnosis of chronic rheumatoid arthritis, and for developing no
                                                                                                                                     1283
                                                                                                                                                                       1223
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                                                                                                                                                                                                                                         1103
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Diagnosing rheumatoid arthritis or comparing expression of mRNA transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-308750/32.
P-PSDB; AAB97370.
                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 14-18; 21pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Shiozawa S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                            Sequence 1634
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                                                                                   AAL47186;
                                                                                                   AAL47186 standard;
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                                                                   22-AUG-2002
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                                DR3; rheumatoid arthritis; antirheumatic; antiarthritic;
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                                                                  (first entry)
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Matches:
Conservative:
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Indels:
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Best Local Similarity:
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27-MAR-2001;
30-MAR-2001;
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                                                         AAC91477
                                                                                                                                                                                                                                                                                                                Sequence 1634 BP;
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                        21-MAR-2001
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SHIOZAWA S.
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                                                         standard;
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2001JP-0090546.
2001JP-0099990.
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                       (first entry)
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The present invention relates to the human DR3 gene, which is associat with rheumatoid arthritis. Certain mutations in the gene can be linked the disease. The sequences can be used to evaluate disease onset and i possibility and to provide therapy and remedies. The present sequence a coding sequence described in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Genomes, particularly DR3 genomic DNA, participating arthritis via mutation, useful in evaluating disease possibility and providing therapy and remedies -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 66-69; 84pp; Japanese.
                                                           CGCATGGGGCTGGACGGCTGCGTGGAAGACTTG
                 ArgMetGlyLeuAspGlyCysValGluAspLeu
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/product= "AAO17879"
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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71
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29-OCT-1999;
30-NOV-1999;
01-DEC-1999;
09-DEC-1999;
20-DEC-1999;
20-DEC-1999;
05-JAN-2000;
06-JAN-2000;
11-FEB-2000;
                                                                                                                    18-FEB-2000;
18-FEB-2000;
22-FEB-2000;
24-FEB-2000;
15-MAR-2000;
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20-JUL-1999
20-JUL-1999
28-JUL-1999
01-SEP-1999
15-SEP-1999
15-SEP-1999
                                                                                    20-MAR-2000;
21-MAR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antirheumatic; cardiant; antianaemic; immunosuppressive;
antidiabetic; nootropic; neuroprotective; hepatotropic; vantiallergic; antiasthmatic; immune related disorder;
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GENENTECH
                              99US-0144732

99US-0146723

99US-0146222

99WO-US20111

99WO-US21547

99US-0162506

99WO-US28313

99WO-US2834341

2000WO-US2834341

2000WO-US284341

2000WO-US284341

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2000WO-US28439

2000WO-US213705

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       disease; autoimmune disease; allergy; ss.
INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           dermatological; antiarthritic;
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anaemia, autoimmune thrombocytopaenia, thyroiditis, diabetes mellitus, immune-mediated renal disease, demyelinating diseases of the central and peripheral nervous systems (such as multiple sclerosis, idiopathic demyelinating polyneuropathy or Guillain-Barre syndrome, and chronic inflammatory demyelinating polyneuropathy), hepatobiliary diseases (such as infectious, autoimmune chronic active hepatitis, primary biliary cirrhosis, granulomatous hepatitis and sclerosing cholangitis), inflammatory bowel disease, gluten-sensitive enteropathy and Whipple's disease, autoimmune or immune-mediated skin disease, (such as bullous disease, autoimmune or immune-mediated skin disease, (such as bullous disorders such as systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis, juvenile chronic arthritis, spondyloarthropathies, systemic sclerosis, idiopathic inflammatory myopathies, Sjogren's systemic systemic vasculitis, sarcoidosis, autoimmune haemolytic syndrome, systemic vasculitis, sarcoidosis, autoimmune haemolytic Thirty three nucleic acids encoding PRO I in the diagnosis and treatment of immune systemic lupus erythematosus, rheumatoid thyroiditis and diabetes mellitus -The present sequence is one of thirty three nucleic acids encoding polypeptides. The PRO polypeptides, anti-PRO antibodies, agonists an antagonists are useful for treating and diagnosing immune related Claim 48; Fig 33 218pp; English. encoding PRO polypeptides which are useful ment of immune related disorders, e.g. us, rheumatoid arthritis, osteoarthritis, encoding

skin

diseases,

multiforme,

contact

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ABK40265
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22-JUN-1999
20-JUL-1999
26-JUL-1999
28-JUL-1999
17-AUG-1999
31-AUG-1999
01-SEP-1999
01-SEP-1999
01-DEC-1999
01-DEC-1999
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11-MAR-1999;
11-MAY-1999;
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                                                                                                                                                           11-FEB-2000;
                                                                                                                                                                                               WO200153486-A1
                                                                                                                                                                                                                                                                               cDNA encoding human
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                                                                                                     22-JUN-1999;
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                                                                                                                                                                                                                                            inflammatory disorder;
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                                                                                                                                                                                                                                                                                                                                                                                                            therapy;
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                                                                                                                                                                                                                                                                                                                                      standard;
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99US-140650P.
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99US-145698P.
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99US-146222P.
99US-151689P.
99US-151689P.
99WO-US20111.
99WO-US20113.
99WO-US28634.
                                                                                                                                                                                                                                 cytostatic; neuroprotective;
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99WO-US12252.
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                                                                                                                                                                                                                                           tumour; malignant tumour; lymphol disorder; stromal disorder; blader; immune disorder; angiogenic
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                                                                                                                                                                                                                                  gene; ss.
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er; blastocoelic disc
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                                                                                                                                                                                                                                           disorder;
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Wood

WI;

AJ, Baker Henzel W,

KP, Chan Kabakoff

ж<sub>С</sub>,

Goddard A, Shelton DL,

Godowski PJ, Tumas À

Watanabe CK;

Gurney

2001-025253/03.

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Best Local S. Query Match: DB:
RESULT 10
AAT89426
ID AAT89
XX AAT89
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XX D2-MA
XX Death
XX Death
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          polypeptides and the polynucleotide sequences encoding them. The PRO polypeptides, agonists, antagonists or anti-PRO antibodies are useful for treating benign or malignant tumours (e.g. renal, kidney, bladder, breast, etc), leukaemias and lymphoid malignancies, other disorders such as neuronal, glial, astrocytal, hypothalamic, glandular, macrophagal, stromal and blastocoelic disorders, inflammatory, immune and angiogenic disorders. The polynucleotide sequences are also useful in gene therapy, ABK40254-ABK40288 encode for the human PRO
                                                                                                                                                                                                                                                                                                                                                                                                                1117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Thirty five nucleic acids encoding PRO polypeptides, useful for treating benign or malignant tumours, leukaemias and lymphoid malignancies, inflammatory, anglogenic and immunologic disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 50;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Marsters SA, Watanabe CK,
                                                                  inflammation;
                                                                              Death domain
                                                                                                         Death domain
                                                                                                                                    02-MAR-1998
                                                                                                                                                              AAT89426;
                                                                                                                                                                                         AAT89426 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 1662 BP;
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                                                                                                                                                                                                                                                                                                               LeuLysArgTrpArgGlnGlnGlnProAlaGlyLeuGlyAlaValTyrAlaAlaLeuGlu
                                       sapiens
                                                                                                                                                                                                                                             CGCATGGGGCTGGACGGCTGCGTGGAAGACTTG 1329
                                                                                                                                                                                                                                                            ArgMetGlyLeuAspGlyCysValGluAspLeu 71
                                                                                                                                                                                                                                                                                                   CTCAAGCGCTGGCGCCAGCAGCCCGCGGGCCTTCGGAGCCGTTTACGCGGCCCTGGAG
                                                                                                                                                                                                                                                                                                                                                       GCAGAGATCGAAGCCGTGGAGGTGGAGATCGGCCGCTTCCGAGACCAGCAGTACGAGATG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fig 23;
                                                               containing 
NF-kappaB;
                                                                                                       containing receptor DR3-V1 cDNA.
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Wood WI;
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Pan J, Pitti RM,
Location/Qualifiers
198..1484
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                                                                 receptor; DR3-V1; ds.
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i RM, Roy MA,
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Smith
                                                                            human; apoptosis;
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Stone DM;
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                                                                                                                                                                                                                                                                                                                                                                                                                domain containing receptor DR3-V1 (see AAW31516), a novel member of the tumour necrosis factor receptor family. It was isolated from a cDNA library derived from cells of a human testis tumour. Related death domain containing receptor DR3 cDNA (see AAT89427) was isolated from a HUVEC cDNA library. The genes have also been identified in cDNA libraries of foetal liver, foetal brain, tonsil and leukocyte. Nucleic acids encoding full-length or mature DR3-V1 can be used to produce recombinant polypeptides in transformed host cells. These polypeptides can be used to treat diseases and disorders associated with the inhibition of apoptosis. Antagonists, such as antibodies raised against DR3-V1, can be used to treat diseases and disorders associated with increased apoptosis and for treating inflammatory
                                       1425
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                                                                                                                                                                                           mat_peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Death domain containing receptor polypeptide(s) DR3 and DR3-V1 for activation of apoptosis and NF-kappaB, antagonists can be u to treat inflammatory diseases
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P-PSDB; AAW31516.
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                                                    ArgMetGlyLeuAspGlyCysValGluAspLeu
                                                                                                      LeuLysArgTrpArgGlnGlnGlnProAlaGlyLeuGlyAlaValTyrAlaAlaLeuGlu
                                                                                                                                                      AlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGluMet
                                       CGCATGGGGCTGGACGGCTGCGTGGAAGACTTG
                                                                                        CTCAAGCGCTGGCGCCAGCAGCAGCCCGCGCGCCTCGGAGCCGTTTACGCGGCCCTGGAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               UNIV MICHIGAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HUMAN GENOME SCI INC
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                                                                                                                                                                                                Alignment
Pred. No.:
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                                                                                                                                                                                                                                                          The invention provides nucleotide sequences encoding death domain containing receptor polypeptides DR3-V1 and DR3. The DR3-V1 cDNA clone is contained in ATCC deposition No. 97456 and DR3 cDNA clone is contained in ATCC deposition No. 97757. Recombinant vectors comprising the nucleic acid sequences and optionally the leader sequences are used for the recombinant production of the proteins. The present sequence represents a cDNA encoding the death domain containing
                                                                             1245
                                                                                                                                                                                                                               Sequence 1783
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                                                                                                                                                                                                                                                                                                                                                                     optionally comprising leader sequence
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17-OCT-1996;
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                                                                                                                                                                                                                                                                                                                                                                                New death domain containing receptor and recombinant vector
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(UNMI) UNIV MICHIGAN.
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                                                                                                \vdash
                                                                                    MetAspAlaValProAlaArgArgTrpLysGluPheValArgThrLeuGlyLeuArgGlu
LeuLysArgTrpArgGlnGlnGlnProAlaGlyLeuGlyAlaValTyrAlaAlaLeuGlu
                                 GCAGAGATCGAAGCCGTGGAGGTGGAGATCGGCCGCTTCCGAGACCAGCAGTACGAGATG
                                           AlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspClnGlnTyrGluMet
                                                                           ATGGACGCGGTCCCAGCGCGCGCGCTGGAAGGAGTTCGTGCGCACGCTGGGGGCTGCGCGAG
                                                                                                                                                                                                                                                                                                                                                                                                                1999-124390/11.
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                                                                                                                                                                                                                                                                                                                                                 Fig 1,
                                                                                                                                                                                                                                                    polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     containing receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                               B₽;
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96US-0013285.
96US-0028711.
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301..1481
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198..300
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198..1484
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368.00
100.00%
100.00%
100.00%
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Matches:
Conservative:
Mismatches:
Indels:
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MetAspAlaValProAlaArgArgTrpLysGluPheValArgThrLeuGlyLeuArgGlu

20

x AAC68776 (1-1783)

US-09-993-234-6\_COPY\_339\_409 (1-71)

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Alignment :
Pred. No.:
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AAC68776
             Percent Similarity:
Best Local Similarity:
Query Match:
   DB:
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                                                                                                                             The present invention provides the protein and coding sequences for two death domain containing receptors, designated DR3 and DR3-V1. These receptors are involved in apoptosis, and the sequences given can be used in the treatment of cancers, infections, cardiovascular disorders such as arrhythmias, ischaemia, aneurysms, arterial occlusive diseases, embolisms and congenital heart defects, neurodegenerative diseases unch as multiple Alzheimer's and Parkinson's diseases, autoimmune disease such as multiple sclerosis, arthritis, diabetes, Graves' disease, asthma and psoriasis,
                                                                                                                                                                                                                                                            Treating graft-versus-host disease, cancer, immunodeficiency autoimmune disease comprising administering an antibody to De Containing Receptor proteins and a second therapeutic agent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1425
                                                                                                                                                                                                                                                                                                                                             Yu G,
                                                                                              Sequence 1783
                                                                                                                                                                                                                                                                                                                                                                                                                                                               22-APR-1999;
28-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; death domain (
autoimmune disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human death domain containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAC68776 standard;
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                                                                                                                                                                                                                                      Example 1; Fig 1; 273pp; English.
                                                                                                                                                                                                                                                                                                            P-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                            (GENT/)
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                                                                       Scores:
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DB; AAB36264.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          domain containing receptor; DR3-V1;
isorder; inflammation; cardiovascula;
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                                                                                             BP;
                                                                                                                    angiogenesis and
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99US-0136741
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                                                                                             564 G;
                                                                                                                    wound healing
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            Length:
Matches:
Conservative:
Mismatches:
Indels:
  Gaps:
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1783
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                                                                                                                                                                embolisms
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RESULT 13
AAV28700
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Percent Similarity:
Best Local Similarity:
                   Alignment Scores: Pred. No.: Score:
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                                                                      This sequence encodes the human apoptosis inducing receptor (AIR) of the invention. AIR is a Type I transmembrane protein, soluble forms of which can be used to regulate cell death in a therapeutic setting. Soluble AIR can also be used in vitro to block apoptosis or AIR-expressing cells, or to screen agonists or antagonists of AIR activity. The cytoplasmic domain of AIR can be used to develop assays for inhibitors of AIR-induced cell death, which is useful to regulate cell death in a therapeutic setting as well as in vitro. Agonists of AIR activity can be used to kill tumour cells that express AIR, or T cells expressing AIR in autoimmune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1245
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                                                                                                                                                                                                                                                                                                       03-OCT-1997;
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                                                                                                                                                                                                                    P-PSDB;
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                                                                                                                                                                                                                                                                                                                           09-APR-1998
                                                                                                                                                                                                                                                                                                                                             WO9814565-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human apoptosis inducing receptor coding sequence.
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                                                         Sequence 1847 BP;
                                                                                                                                                                       Claim
                                                                                                                                                                                       DNA encoding apoptosis inducing receptor - which is Type transmembrane protein, useful for regulating cell death
                                                                                                                                                                                                                                                 Perkins PA;
                                                                                                                                                                                                                                                                  (IMMV ) IMMUNEX CORP.
                                                                                                                                                                                                                                                                                                                                                                                                              Homo
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DB; AAW57045.
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                                                                                                                                                                     2; Page 28-30; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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236..1489
/*tag= a
                                                                                                                                                                                                                                                                                                                                                               /product= AIR
                                                        316 A; 605 C;
3.63e-40
368.00
100.00%
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                                                        579 G;
Length:
Matches:
Conservative:
Mismatches:
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                                                        347 T;
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                                                                                                                                                                                                                                                                                                                                                                                                                                       human; cell death regulator; death; autoimmune disease;
                                                        0 other;
1847
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; DR3; rheumatoid arthritis; antirheumatic; antiarthritic;
gene therapy; gene; ds.
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                                     intron
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                                                                                                                                                                                                                                                                                                                                                                                                           Human DR3 gene associated with rheumatoid arthritis
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1477..1644
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128..635
        3069..312
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                                 1822..3068
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                                                               743..182
                                                                         'number-
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Alignment
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                                                         US-09-993-234-6_COPY_339_409 (1-71) x
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Best Local Similarity:
Query Match:
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27-MAR-2001;
30-MAR-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        exon
                                                                                                                   No.:
                                                                                                                                                           The present invention relates to the human DR3 gene, which is associated with rheumatoid arthritis. Certain mutations in the gene can be linked to the disease. The sequences can be used to evaluate disease onset and its possibility and to provide therapy and remedies. The present sequence is the gene of the invention.
                                                                                                                                                                                                                          Genomes, particularly DR3 genomic DNA, participating arthritis via mutation, useful in evaluating disease possibility and providing therapy and remedies -
                                                                                                                                                                                                                                                                                                                                                                                                                       exon
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                                                                                                                                          Sequence 4825
                                                                                                                                                                                                            Claim 1; Page 64-66; 84pp; Japanese.
                                                                                                                                                                                                                                                                                           (NEWI-)
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                        AlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGluMet
                                                                                                                           Scores:
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                                                                                                                                                                                                                                                                                            NEW IND RES ORG. SHIOZAWA S.
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; 2001JP-0090546.
; 2001JP-0099990.
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4434..4825
/*tag= t
/numb
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/number= 7
3530..3577
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3334..3529
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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cytostatic; gene; ds.
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Barash SC, Ruben MS

2001-565190/63.

Nucleic used in such as acid encoding novel connective tissue associated polypeptides, diagnosing, preventing, treating or ameliorating a disorder cancer or rheumatoid arthritis -

Disclosure; SEQ ij ĕ 1577; 673pp; English.

The present invention relates to the isolation of novel human connective tissue related polypeptides (AROU86435-AAU86923) and the polynucleotide (cDNA and genomic) sequences encoding them. The sequences of the invention are useful in the diagnosis, treatment, prevention and/or prognosis of diseases associated with connective tissue(s), including cancer. The polynucleotide sequences of the invention are also useful in gene therapy. ABK42102-ABK43116 represent genomic sequences encoding the novel human connective tissue related polypeptides. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO ftp.wipo.int/pub/published\_pct\_sequences

Sequence 10797 BP; 2003 A; 3215 Ç 3378 <u>ດ</u> 2201 Ŧ; 0 other;

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Best Local Similarity:
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CGCATGGGGCTGGACGGCTGCGTGGAAGACTTG

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Search completed: April 6, 2003, 16:53:36 Job time : 75.109 secs

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## ALIGNMENTS

RESULT 1 US-08-815-469-3

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Description US/08815469 Patent No. 6153402
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                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: NO. (
FILING DATE: 06-FEB-1997
PRIOR APPLICATION DATA:
                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Yu, Guo-Liang
APPLICANT: Ni, Jian
APPLICANT: Dixit, Vishva
                  PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Dillon, Patrick TITLE OF INVENTION: Death I
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                                  APPLICATION NUMBER: US 60/028,711 FILING DATE: 17-OCT-1996
                                                                                                                                            CLASSIFICATION:
                                                                                                                                                             APPLICATION NUMBER: US/08/815,469 FILING DATE: HEREWITH
                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Sterne, Kessler, STREET: 1100 New York Ave.,
                                                                                                                                                                                                                                                                                                               20005-3934
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Gentz, Reiner L.
NUMBER:
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US 60/013,285
                                                                                                          6153402 Yet Assigned
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Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Steffe, Eric n. 6,688
REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 1488
TELECOMMUNICATION INFORMATION:
"ERIEPHONE: 202-371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 3: SEQUENCE CHARACTERISTICS:
                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb:
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
                                           CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/928,069
FILING DATE: 11-Sep-1997
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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APPLICANT: Ashkenazi, Avi J.
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                             CLASSIFICATION:
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                                                                                                                                                            ZIP: 94080
                                                                                                                                                                        COUNTRY:
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STRANDEDNESS: doub
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Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 9, Application US/08828683A
Patent NO. 6469144
GENERAL INFORMATION:
APPLICANT: AShkenazi, Avi J.
TITLE OF INVENTION: Apo-2 LI i
NUMBER OF SEQUENCES: 28
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INFORMATION FOR SEQ ID NO:
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LENGTH: 1634 base pair
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TYPE: Nucleic Acid
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REGISTRATION NUMBER:
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MEDIUM TYPE: 3.5 inch, 1.44 Mb |
COMPUTER: IBM.PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/625328
FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
                                                                                                                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/828,683A
FILING DATE: 31-Mar-1997
CLASSIFICATION: <Unknown>
                 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
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ZIP: 94080
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ER: P1052R1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1, Application Patent No. 6153402
                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICATION NUMBER: US/08/
FILING DATE: HEREWITH
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: NO. 61
FILING DATE: 06-FEB-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/
FILING DATE: 17-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
APPLICANT:
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MEDIUM TYPE: Floppy disk
                                                                                                                                    SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
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                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                         T: Dixit, Vishva
T: Gentz, Reiner L.
T: Dillon, Patrick J.
INVENTION: Death Domain Containing Receptors
F SEQUENCES: 17
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TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                           E: Sterne, Kessler,
1100 New York Ave.,
                                                                                                                                                                                                                                                                                                                                                                                                   Yu, Jian
Ni, Jian
''t, V
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                60/028,711
                                                            6153402 Yet Assigned
                                                                                                                                                                                                                                                                                             Goldstein & Fox, P.L.L.C NW, Suite 600
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Matches:
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                                                                                                                                                                                                                                                                                    Sequence 10, AFF--
Sequence 10, AFF--
Sequence 10, AFF---
                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Beutler,
APPLICANT: Bazzoni,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 202-371-2540 INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS:
                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                      1365
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APPLICATION NUMBER: US 6
FILING DATE: 12-MAR-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                               1425 CGCATGGGGCTGGACGGCTGCGTGGAAGACTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1245 ATGGACGCGGTCCCAGCGCGCGCGCGCAAGGAGTTCGTGCGCACGCTGGGGCTGCGCGAG
                                                                                                                                                                                                                        TITLE OF INVENTION:
                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                         61
                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY:
                                                                                                            COUNTRY: USA
ZIP: 77210-4433
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                                                                                                                                       STATE:
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             APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                        Application US/08762308
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P.O. Box 4
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K 4433
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CLASSIFICATION: 530 PRIOR APPLICATION DATA:

APPLICATION NUMBER: FILING DATE: 05-AP

US 08/224,593

VQ.

1548

1608

57 AlaAlaLeuGluArgMetGlyLeuAspGlyCysValGluAspLeu 71

CGCGTGCTCAGGGACATGGACCTGCTGGGTTGCCTGGAAAACATA 1652

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1488 CACGAGATCGAGCGGCTGGAGCTGGAGAACGGGCGCCACCTGCGCGAGGCGCAGTACAGC 1547

21 AlaGluIleGluAlaValGluIleGlyArg---PheArgAspGlnGlnTyrGlu 39

40 MetLeuLysArgTrpArgGlnGlnGlnPro-----AlaGlyLeuGlyAlaValTyr 56

ATGCTGGCGGCCTGGCGCGCGCGCACGCCGCGCGAGGCCACGCTGGAGCTGCTGGGC 1607

1428 GTGGACGGCGTGCCCCCGTCGCGCTGGAAGGAGTTGGTGCGGCGGCTGGGACTGAGCGAG 1487

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Query Match:

Percent Similarity: Best Local Similarity:

1.21e-12 156.00 61.33% 48.00%

Conservative: Mismatches: Indels:

Gaps:

Length: Matches:

2440 36 10 25

Pred. No.: Score:

US-09-993-234-6\_COPY\_339\_409 (1-71) x US-09-513-007-1 (1-2440)

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Alignment Scores
                                  ; NAME/KEY: CDS
; LOCATION: (29)
US-09-513-007-1
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                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1, Applic Patent No. 6406907
                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/122,156
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 8
                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/513,007
CURRENT FILING DATE: 2000-02-25
                                                                                                                                                                                                                                                                                         APPLICANT: Mwangi, Simon
TITLE OF INVENTION: BOVINE TUMOR NECROSIS FACTOR RECEPTOR-1
TITLE OF INVENTION: AND METHODS OF USE
                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Taylor, J. Michael APPLICANT: Kehrli, Jr., Marcus
                                                                                                                                                                                                                                                                                  FILE REFERENCE: 08411-018001
                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                             TYPE: DNA
ORGANISM: Bos taurus
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                                                                                        FEATURE:
                                                                                                                                          LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Kitchell, Barbara S.
REGISTRATION NUMBER: 33,928
REFERENCE/DOCKET NUMBER: UT
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1362 CTCGTGCTTTCCAAGATGAACCTGGCTGGGTGCCTGGAGAATATC 1406
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STRANDEDNESS: single
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                                                 (294)...(1706)
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Indels:
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; SEQ ID NO 236
; LENGTH: 427
; TYPE: DNA
; ORGANISM: Homo sapien
US-09-397-787-236
                                                                                                                                                                                                                                                                                                                              US-09-993-234-6_COPY_339_409 (1-71) x US-09-397-787-236 (1-427)
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Mitcham, Jennifer L.
APPLICANT: King, Gordon E.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR OVARIAN
TITLE OF INVENTION: CANCER THERAPY AND DIAGNOSIS
FILE REFERENCE: 210121.46662
CURRENT APPLICATION NUMBER: US/09/397,787
CURRENT FILING DATE: 1999-09-16
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236 CGCGTGCTCCGCGACATGGACCTGCTGGGCTGCCTGGAGGACATC 280
                                                                                   176
                                                                                                                                                            57 AlaAlaLeuGluArgMetGlyLeuAspGlyCysValGluAspLeu 71
                                                                                                                 40 MetLeuLysArgTrpArgGlnGlnGlnPro-----AlaGlyLeuGlyAlaValTyr 56
                                                                                                                                                                                                21 AlaGluIleGluAlaValGluValGluIleGlyArg---PheArgAspGlnGlnTyrGlu 39
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                                                                                                                                                                                                                                                                                 1 MetAspAlaValProAlaArgArgTrpLysGluPheValArgThrLeuGlyLeuArgGlu 20
                                                                             ATGCTGGCGACCTGGAGGCGGCGCACGCCGCGGGGGCGAGGCCACGCTGGAGCTGCTGGGA 235
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US-08-050-319B-24

Sequence

24, Application US/08050319B 5. 5633145

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; LOCATION:
US-08-050-319B-24
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INFORMATION FOR SEQ
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 617-8999
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MEDIUM TYPE: Floppy disk
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APPLICANT: M.J.C. Turner, F.M Brennan
TITLE OF INVENTION: Modified human TNFalpha (Tumor
TITLE OF INVENTION: Modified human TNFalpha) Receptor
                                                                                                           1361
                                                                                                                                                                                                                                      1241
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CURRENT APPLICATION DATA:
                                            1421 CGCGTGCTCCGCGACATGGACCTGCTGGGCTGCCTGGAGGACATC 1465
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                                                                                                                                                                                         21 AlaGluIleGluAlaValGluValGluIleGlyArg---PheArgAspGlnGlnTyrGlu 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: doub
TOPOLOGY: linear
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                                                                                                                                                                                                                                    GTGGAGAACGTGCCCCCGTTGCGCTGGAAGGAATTCGTGCGGCGCCTAGGGCTGAGCGAC 1300
                                                                     AlaAlaLeuGluArgMetGlyLeuAspGlyCysValGluAspLeu 71
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ID NO: 2
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RESULT 10
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US-08-465-982-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (415) 327-32 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/465,982
                                                                                                       1361
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TELECOMMUNICATION INFORMATION:
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MEDIUM TYPE: Floppy disk
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APPLICANT: M.J.C. Turner, F.M Brennan
TITLE OF INVENTION: Modified human TNFalpha (Tumor
TITLE OF INVENTION: Necrosis Factor alpha) Receptor
                                1421 CGCGTGCTCCGCGACATGGACCTGCTGGGCTGCCTGGAGGACATC 1465
                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                  1241 GTGGAGAACGTGCCCCCGTTGCGCTGGAAGGAATTCGTGCGGCGCCCTAGGGCTGAGCGAC 1300
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                                                                                                                                                                                                    21 AlaGluIleGluAlaValGluValGluIleGlyArg---PheArgAspGlnGlnTyrGlu 39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
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                                                       AlaAlaLeuGluArgMetGlyLeuAspGlyCysValGluAspLeu 71
                                                                                                     ATGCTGGCGACCTGGAGGCGGGGCACGCCGCGGGGGCGAGGCCACGCTGGAGCTGCTGGGA 1420
                                                                                                                                     MetLeuLysArgTrpArgGlnGlnGlnPro-----AlaGlyLeuGlyAlaValTyr 56
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               GENERAL INFORMATION:
APPLICANT: Rosen, C
APPLICANT: Kao, Pet
                                                                   Sequence 3, Application US/09505250A Patent No. 6329148
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   TITLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (760) 603-3820 INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Laurel Spear Bernstein REGISTRATION NUMBER: 37,280 REGISTRATION NUMBER: RTS-TELECOMMUNICATION INFORMATION: TELEPHONE: (760) 931-9200
                                                                                                                                                                                                                                                                                                                                                            1342 GTGGAGAACGTGCCCCCGTTGCGCTGGAAGGAATTCGTGCGGCGCCTAGGGCTGAGCGAC 1401
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                                                                                                                                                                                                                              1462
                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
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FILING DATE: June 26, 1998
CLASSIFICATION: 514
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   OF INVENTION:
                                                                                                                                                        CGCGTGCTCCGCGACATGGACCTGCTGGGCTGCCTGGAGGACATC 1566
                                                                                                                                                                                        AlaAlaLeuGluArgMetGlyLeuAspGlyCysValGluAspLeu 71
                                                                                                                                                                                                                        ATGCTGGCGACCTGGAGGCGGCGCACGCCGCGCGCGCGAGGCCACGCTGGAGCTGCTGGGA 1521
                                                                                                                                                                                                                                                          MetLeuLysArgTrpArgGlnGlnGlnPro-----AlaGlyLeuGlyAlaValTyr 56
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Microsoft Word 97
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Synergistic Anti-Cancer Therapy with
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Percent Similarity:
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Query Match:
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; LOCATION: (25)
US-09-505-250-3
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CURRENT FILING DATE: 2000-02-15
NUMBER OF SEQ ID NOS: 4
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 3
LENGTH: 2161
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Patent No.
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 1L 107268
FILING DATE: 12-OCT-1993
                                                                                                                                                                                                                                                                                                                                                           APPLICANT: BATKIN, Michael TITLE OF INVENTION: MOLECU TITLE OF INVENTION: THE TN
                                                                                                      CURRENT APPLICATION DATA:
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ADDRESSEE: BROWDY AND NEIMARK
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                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
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                                                                                                                      MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
                                                                                                                                                                                                                                                   CITY: Washington STATE: D.C.
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                                                                        FILING DATE:
                                                                                       APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                      419 Seventh Street, N.W.,
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                                                                                       US/08/321,668
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TELEPHONE: 202-628-5197

NAME: BROWDY, Roger L.
REGISTRATION NUMBER: 25
REFERENCE/DOCKET NUMBER:

25,618

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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                             Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1,
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TELEX: 248633
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1402
                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1342 GTGGAGAACGTGCCCCCGTTGCGCTGGAAGGAATTCGTGCGGCGCCCTAGGGCCTGAGCGAC 1401
                                                                                                               MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: BROWDY AND NEIMARK
                                                                                                                                                                                                                                                                                     APPLICANT: BRAKEBUSCH, Cord
APPLICANT: VARROLDMEEV, Eugene
APPLICANT: BATKIN, Michael
TITLE OF INVENTION: MOLECULES J
TITLE OF INVENTION: THE TNF REC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1462
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LOCATION:
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            APPLICATION NUMBER: US/08/837,941 FILING DATE: 28-APR-1997
                                                                                                                                                                                                                     STREET:
CLASSIFICATION:
                                                                                                                                                                  COUNTRY:
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CGCGTGCTCCGCGACATGGACCTGCTGGGCTGCCTGGAGGACATC 1566
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                                                                                                                                                  20004
                                                                                                                                                                                    Washington : D.C.
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                                                                                                                 Floppy disk
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151.00
62.67%
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                                                                                                                                                                                                                                                                                                                                                                            David
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Matches:
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                                                               Version #1.30
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RESULT 14
US-08-126-016-1
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Best Local Similarity:
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US-08-837-941-1
                                                                                                                                                                                                                                                                         Sequence 1 Patent No.
                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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                                                                                                                                                                                              APPLICANT:
APPLICANT:
                                                                                                                              APPLICANT: BRAKEBUSCH, CORD
APPLICANT: ADERKA, DAN
TITLE OF INVENTION: EXPRESSION OF THE RECOMBINANT TUMOR
TITLE OF INVENTION: NECROSIS FACTOR BINDING PROTEIN I (7)
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NAME/KEY:
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ATTORNEY/AGENT INFORMATION:
NAME: BROWDY, Roger L.
REGISTRATION NUMBER: 25,6
COMPUTER READABLE FORM:
                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Browdy an
                                                                                                                    NUMBER OF SEQUENCES:
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TELEFAX: 248633
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                                            CITY: Washington
STATE: DC
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                           COUNTRY:
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KEMPER, OLIVER
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                                                                         Street, N.W.,
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Matches:
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SOFTWARE: Patentin Rel CURRENT APPLICATION DATA:

APPLICATION NUMBER:

US/08/126,016

Release #1.0, Version #1.25

COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS

MEDIUM TYPE:

Floppy disk

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Percent Similarity:
Best Local Similarity:
Query Match:
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US-08-054-970-1
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                                                                                                                                                    Sequence 1, Application US/08054970 Patent No. 6395267
                                                                                                    GENERAL INFORMATION:
APPLICANT: WALLACH, David
APPLICANT: BRAKEBUSCH, Cord
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 24-SEP-1993
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/625668
FILING DATE: 13-DEC-1990
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LENGTH: 2175 base pairs
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TELEFAX: 202-737-3528
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                                                        NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
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                                                                                     TITLE OF INVENTION:
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LOCATION:
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REGISTRATION NUMBER: 25,618
REFERENCE/DOCKET NUMBER: WA
STREET: 412
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                                                                                                                                                                                                                                                                                                                       40 MetLeuLysArgTrpArgGlnGlnGlnPro-----AlaGlyLeuGlyAlaValTyr 56
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                                           BROWDY AND NEIMARK
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Matches:
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Percent Similarity:
Best Local Similarity:
Query Match:
Search completed: April Job time: 17.9856 secs
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US-08-054-970-1
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENCTH: 2175 bass Types
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APPLICATION NUMBER: US,
FILING DATE: 03-MAY-19
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
TELEPAX: 202-737-3528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Townsend, Kevin G.
REGISTRATION NUMBER: 34,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
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                                                             1522 CGCGTGCTCCGCGACATGGACCTGCTGGGCTGCCTGGAGGACATC 1566
                                                                                                                           1462 ATGCTGGCGACCTGGAGGCGGCGCGCCGCGCGCGCGAGGCCACGCTGGAGCTGCTGGGA 1521
                                                                                                                                                                                             MOLECULE TYPE:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                    21 AlaGluIleGluAlaValGluValGluIleGlyArg---PheArgAspGlnGlnTyrGlu 39
                                                                                            57 AlaAlaLeuGluArgMetGlyLeuAspGlyCysValGluAspLeu 71
                                                                                                                                                         40 MetLeuLysArgTrpArgGlnGlnGlnPro------AlaGlyLeuGlyAlaValTyr 56
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TOPOLOGY: 111
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Conservative:
Mismatches:
Indels:
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Result
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-Q-/cgn2_1/USPTO_spool/US09993234/runat_27032003_115459_15466/app_query.fasta_1.2346
-DB-Published_Applications_NA -OFNT-fastap -SUFFIX-rnpb -MINMATCH-0.1
-LOOPCL-0 -LOOPEXT-0 -UNITS-bits -START-1 -END-1 -MATRIX-blosum62
-TRANS-human40.cdi -LIST-45 -DOCALIGN-200 -THR_SCORE-pct -THR_MAX-100
-THR_MIN-0 -ALIGN-15 -MODE-LOCAL -OUTFMT-pto -NORM-ext -HEAPSIZE-500 -MINLEN-0
-MAXLEN-200000000 -USER-US0993234_eCGN_1_1_249_erunat_2703203_115459_15466
-NCPU-6 -ICPU-3 -NO_XLPXY -NO_MAP -LARGEQUERY -NEG_SCORES-0 -WAIT -LONGLOG
-DEV_TIMEOUT-120 -WARN_TIMEOUT-30 -THREADS-1 -XGAPOP-10 -XGAPOP-10 -SGAPOP-6
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Perfect score:
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                                                                                                                                                                 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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                                                                                    Score
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length: 2000000000
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Ygapop 10.0 , X
Fgapop 6.0 , I
Delop 6.0 , I
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Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                  /cgn2_6/ptodata/1/pubpna/US07_NEW_PUB_seq:*
/cgn2_6/ptodata/1/pubpna/PCTUS_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US08_NEW_PUB_seq:*
/cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
                                                                                                                                                                                                                               /cgn2_6/ptodata/1/pubpna/US09_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US60_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US60_PUBCOMB.seq:*
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   1254
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US-10-112-793-9
US-10-112-193-11
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                                                                                                                                     SUMMARIES
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               Sequence 3,
Sequence 9, A
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Sequence 9, 1
Sequence 9, 1
Sequence 11,
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US-08-916-625B-1 US-09-811-088-5 US-09-757-421-3 US-09-811-088-3 US-09-911-088-3 US-09-9757-421-1 US-09-991-496-5 US-09-991-496-99 US-09-991-496-99	09-792-356- -09-899-422- -10-120-397 08-916-625B 10-076-773- -10-067-615- -09-874-138 -10-005-842- 09-992-964- -09-887-879 -10-052-798	-09-993 -09-333 -09-333 -09-764 -09-754 -09-899-7 -09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-899-09-99-899-09-899-09-99-899-09-99-899-09-99-899-09-99-99-99-99-99-99-99-99-99-99-99-9
1, 15 3, 1 3, 1 5, 1 99,	16, e 1, 3, 3, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,	e 1, 14, 10, 14, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16

## ALIGNMENTS

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RESULT 1
US-09-333-966-3
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                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                   APPLICANT:
APPLICANT:
                                          COMPUTER READABLE FORM:
MEDIUM TYPE: F10ppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                         APPLICANT: Dillon, Patrick J.
TITLE OF INVENTION: Death Domain Containing Receptors
NUMBER OF SEQUENCES: 17
                               CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                            CITY: Washington STATE: DC
                                                                                                                                COUNTRY: USA
ZIP: 20005-3934
                                                                                                                                                                                                   STREET:
FILING DATE:
               APPLICATION NUMBER:
                                                                                                                                                                                                                   ADDRESSEE:
                                                                                                                                                                                               E: Sterne, Kessler,
1100 New York Ave.,
                                                                                                                                                                                                                                                                                                                                     Ni, Jian
                                                                                                                                                                                                                                                                                                   Dixit, Vishva
Gentz, Reiner L.
                                                                                                                                                                                                                                                                                                                                                 Yu, Guo-Liang
                                                                                                                                                                                                                 Kessler,
               us/09/333,966
                                                                                                                                                                                               Goldstein & Fox, P.L.L.C. NW, Suite 600
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RESULT 2
US-10-081
US-10-081
; Sequence 9, Application US/10081280
; Patent No. US20020165157A1
; GENERAL INFORMATION:
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DB:
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INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
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REFERENCE/DOCKET NUMBER: 146
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
                                                                                                                                                                                                                                                                                                                                               1135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 60/028,711 FILING DATE: 17-OCT-1996 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 60/013,285
FILING DATE: 12-MAR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Steffe, Eric K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                   61
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LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                       21 AlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGluMet 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid STRANDEDNESS: doub TOPOLOGY: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ENGTH:
ZIP: 94080
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
                                                                                                           TITLE OF INVENTION: Apo-2 LI AND NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
                                                                                                                                                                           APPLICANT: Ashkenazi, Avi J.
                                                                                                                                                                                                                                                                                                                                            CGCATGGGGCTGGACGGCTGCGTGGAAGACTTG 1227
                                                                                                                                                                                                                                                                                                   ArgMetGlyLeuAspGlyCysValGluAspLeu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATGGACGCGGTCCCAGCGCGCGCGCTGGAAGGAGTTCGTGCGCACGCTGGGGCTGCGCGAG
                                                             STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STATE: California
                                                       COUNTRY:
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Conservative:
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Best Local Similarity:
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INFORMATION FOR SEQ ID NO: 9:
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                                                                                            COUNTRY:
                                                                                                                                          STREET: 1 DNA Way
                                                                                 94080
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US-09-993-234:6_COPY_339_409 (1-71) x US-10-081-280-9 (1-1634)
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                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1103 ATGGACGCGGTCCCCAGCGCGCGCGCTGGAAGGAGTTCGTGCGCACGCTGGGGGCTGCGCGAG 1162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          41 LeuLysArgTrpArgGlnGlnGlnProAlaGlyLeuGlyAlaValTyrAlaAlaLeuGlu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21 AlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGluMet 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 MetAspAlaValProAlaArgArgTrpLysGluPheValArgThrLeuGlyLeuArgGlu 20
                      COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-5416
TELEFAX: 415/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Apo-2 LI AND Apo-3 POLYPEPTIDES NUMBER OF SEQUENCES: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE DESCRIPTION: SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 1634 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/829,270
FILING DATE: 31-Mar-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GCAGAGATCGAAGCCGTGGAGGTGGAGATCGGCCGCTTCCGAGACCAGCAGTACGAGATG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CTCAAGCGCTGGCGCCAGCAGCCGCGCGCGCCTCGGAGCCGTTTACGCGGCCCTGGAG
                                                                                                                                                                  CITY: South San Francisco
STATE: California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007R1
                                                                                                                                                                                                                                           ADDRESSEE: Genentech, Inc
WinPatin (Genentech)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NO:
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CURRENT APPLICATION DATA:

PRIOR APPLICATION DATA:

CLASSIFICATION: <Unknown>

APPLICATION NUMBER: US/10/112,793 FILING DATE: 28-Mar-2002

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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                      US-10-112-193-11
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                                                                                                                                                                                                                                                                               Sequence 11, Application US/10112193 Publication No. US20030004313A1 GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
                                                                                                                                                                                                                        APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-3 POLYPEPTIDE
NUMBER OF SEQUENCES: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Marschang, Diane L.
                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
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APPLICATION NUMBER: 08/625328
FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 1634 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
                                                                                                                 COUNTRY:
                                                                                                                                STATE: California
                                                                                                                                                  CITY: South San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 35,600
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Query Match:
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US-09-993-234-9
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                                                                                                                                                                                                                                                                                                                                              Sequence 9, Application US/09993234 Patent No. US20020146768A1 GENERAL INFORMATION:
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REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1052R1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 60/026943
FILING DATE: 23-Sep-1996
ATTORNEY/AGENT INFORMATION:
                   COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech) CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech,
                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 inch,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                     APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-2 LI AND Apo-3 POLYPEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ArgMetGlyLeuAspGlyCysValGluAspLeu
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FILING DATE: 28-Mar-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                               STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 1634 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
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APPLICATION NUMBER: US/09/993,234
                                                                                                                                                              COUNTRY: USA
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Matches:
Conservative:
Mismatches:
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Best Local Similarity:
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ZIP: 20005-3934

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compartible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Ve
                                                                                                                                                                                                                                                                                                                                                                                Sequence 1, Application US/09333966 Patent No. US20020009773A1
                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                   APPLICANT: Dixit, Vishva
APPLICANT: Gentz, Reiner L.
APPLICANT: Dillon, Patrick J.
TITLE OF INVENTION: Death Dom
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No.:
                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                          STREET: 1100 New CITY: Washington
                                                                                                                                                    STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61 ArgMetGlyLeuAspGlyCysValGluAspLeu 71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             41 LeuLysArgTrpArgGlnGlnGlnProAlaGlyLeuGlyAlaValTyrAlaAlaLeuGlu 60
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                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION: TELEPHONE: 415/225-5416
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE DESCRIPTION: SEQ ID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CTCAAGCGCTGGCGCCAGCAGCCAGCCCGCGGGCCTCGGAGCCGTTTACGCGGCCCTGGAG 1282
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 1634 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007P1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ELEFAX: 415/952-9881
ELEX: 910/371-7168
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Matches:
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                    Version #1.30
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RESULT 7
US-10-092-154-1577
; Sequence 1577, Application US/10092154
; Publication No. US20030054375A1
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Query Match:
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Prior Application removed - See File Wrapper or Palm SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 1577
                                               CURRENT APPLICATION NUMBER: US/10/092,154
CURRENT FILING DATE: 2002-03-07
NUMBER OF SEQ ID NOS: 2003
                                                                                              APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PC009C1
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                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 6(
FILING DATE: 17-OCT-1996
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REGISTON NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 14
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
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ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER:
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TYPE: nucleic acid
STRANDEDNESS: double
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US-09-764-847-1577
; Sequence 1577, Application US/09764847
; Patent No. US20020132787A1
; GENERAL INFORMATION:
APPLICANT: Rosen et al.
; TITLE OF INVENTION: Nucleic Acids, Proteins,
; FILE REFERENCE: PC009
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DB:
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-764-847-1577
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CURRENT FILING DATE: 2001-01-17
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TYPE: DNA
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ArgMetGlyLeuAspGlyCysValGluAspLeu 71
                                                                                 GCAGAGATCGAAGCCGTGGAGGTGGAGATCGGCCGCTTCCGAGACCAGCAGTACGAGATG
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Best Local Similarity:
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CURRENT FILING DATE: 2002-05-02
PRIOR APPLICATION NUMBER: US/60/233,133
PRIOR FILING DATE: 2000-09-18
PRIOR APPLICATION NUMBER: US/60/234,009
PRIOR FILING DATE: 2000-09-20
PRIOR APPLICATION NUMBER: US/60/234,034
PRIOR APPLICATION NUMBER: US/60/234,034
PRIOR APPLICATION NUMBER: US/60/234,034
PRIOR APPLICATION NUMBER: US/60/234,509
PRIOR APPLICATION NUMBER: US/60/234,509
PRIOR FILING DATE: 2000-09-22
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NUMBER OF SEQ ID NOS: 1392

SOFTWARE: PatentIn version 3.0

SEQ ID NO 1389

LENGTH: 1743

TYPE: 1743
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PRIOR

FILING DATE: 2000-11-02
APPLICATION NUMBER: US 60/290,029

APPLICATION NUMBER: US 60/222,880

FILING DATE: 2001-05-11

APPLICATION NUMBER: US 60/222,040 FILING DATE: 2000-07-31

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LENGTH: 2130
                                                     CURRENT APPLICATION NUMBER: US/09/898,234
CURRENT FILING DATE: 2001-07-03
PRIOR APPLICATION NUMBER: 09/525,998
PRIOR FILING DATE: 2000-03-15
                     PRIOR PRIOR
                                                                                                                                                           TITLE OF INVENTION: THE Receptors, TNF Binding Proteins and DNAs Coding TITLE OF INVENTION: Them
                                                                                                                                                                                                    APPLICANT: Hauptmann, Rudolph
APPLICANT: Himmler, Adolph
APPLICANT: Maurer-Pogy, Ingrid
APPLICANT: Stratowa, Christian
                                                                                                                                          FILE REFERENCE: 98,385-
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                 APPLICATION NUMBER: 08/383,676 FILING DATE: 1995-02-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 60/297,457 FILING DATE: 2001-06-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 60/303,459
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NUMBER: 08/153,287
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SEQ ID
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                   SOFTWARE:
                                PRIOR FILING DATE: 199 NUMBER OF SEQ ID NOS:
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                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 08/477,639
PRIOR FILING DATE: 1955-06-07
                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 09/792,356 PRIOR FILING DATE: 2000-02-23
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PRIOR FILING DATE: 1990-04-20
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PRIOR APPLICATION NUMBER: 07/821,750
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                                                                                                                    APPLICATION NUMBER: 08/153,287 FILING DATE: 1993-11-17
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; LOCATION: (245)..(1627)
US-09-899-429A-24
  Pred. No.:
Score:
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US-09-792-356-14
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APPLICANT: Hauptmann, Ruuver
APPLICANT: Himmler, Adolph
APPLICANT: Maurer-Fogy, Ingrid
APPLICANT: Maurer-Fogy, Ingrid
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                                                                                                                                                                                                               SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 14
LENGTH: 2173
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                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/792,356
CURRENT FILING DATE: 2001-08-17
PRIOR APPLICATION NUMBER: 08/477,639
PRIOR FILING DATE: 1995-06-07
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PRIOR FILING DATE: 1993-11-17
PRIOR APPLICATION NUMBER: 07/821,750
PRIOR FILING DATE: 1992-01-02
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PRIOR FILING DATE: 1995-02-01
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ORGANISM: Artificial Sequence
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LOCATION: (245)..(1630)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/899,422
CURRENT FILING DATE: 2001-08-21
PRIOR APPLICATION NUMBER: 09/525,998
PRIOR FILING DATE: 2000-03-15
PRIOR APPLICATION NUMBER: 08/383,676
PRIOR APPLICATION NUMBER: 08/383,676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: THE Receptors, THE Binding Proteins TITLE OF INVENTION: Them FILE REFERENCE: 98,385-H
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE: 1995-02-01
PRIOR APPLICATION NUMBER: 08/153,287
PRIOR FILING DATE: 1993-11-17
PRIOR APPLICATION NUMBER: 07/821,750
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LOCATION: (245)..(1630)
OTHER INFORMATION: Desc
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1352 GTGGATGGCGTGCCTCCGACACGCTGGAAGGAGTTCATGCGGCTCCTGGGGCTGAGCGAG
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Search completed: April Job time: 36.0348 secs
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Best Local Similarity:
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US-09-970-532-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: BOWNE TUMOR NECROSIS FACTOR RECEPTOR-1
TITLE OF INVENTION: AND METHODS OF USE
FILE REFERENCE: 08411-018001
CURRENT APPLICATION NUMBER: US/09/970,532
CURRENT FILING DATE: 2001-10-03
PRIOR APPLICATION NUMBER: 09/513,007
PRIOR PILING DATE: 2000-02-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Taylor, J. Michael
APPLICANT: Kehrli, Jr., Marcus
APPLICANT: Lee, Eun-Kyung
APPLICANT: Mwangi, Simon
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ORGANISM: Bos taurus
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## ALIGNMENTS

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source	FEATURES					COMMENT	JOURNAL		TITLE	AUTHORS	REFERENCE			ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION			DEFINITION	LOCUS	RESULT 1 AW268610	
1632 /organism="Homo sapiens"	High quality sequence stop: 455. Location/Qualifiers.	Possible reversed clone: polyT not found Seq primer: -40UP from Gibco	Possible reversed clone: similarity on wrong strand	This cione is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.	Email: cgapbs-r@mail.nih.gov	Contact: Robert Strausberg, Ph.D.	Unpublished (1997)	Tumor Gene Index	National Cancer Institute, Cancer Genome Anatomy Project (CGAP),	NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.	1 (bases 1 to 632)	Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;	Homo sapiens	human.	EST.	AW268610.1 GI:6655640	AW268610	PRECURSOR ;, mRNA sequence.	IMAGE:2815679 3' similar to SW:WSL1 HUMAN Q93038 WSL-1 PROTEIN	x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clo	AW268610 632 bp mRNA linear EST 03-JAN-2000		

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                         PRECURSOR;, mRNA sequence. A1807913
                                                         AI807913
698 bp mRNA linear EST 19-DEC wf52c03.x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE: 3359204 3' similar to SW:WSL1_HUMAN Q93038 WSL-1 PROTEIN
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/clone_lib="Soares_NFL_T_GBC_S1"
/lab_host="DHIOB"
               GI:5394479
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99.1%;
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                                                                                                                                                                                                                                                                                                                                  CCTTGCACGGAGCCCTGCGGCAACTCCACCTGCCTTGTGTGTCCCCAAGACACCTCTTG
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                                                                                                                                                                                                                                                                                    GCCTGGGAGAACCACCATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGCC
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                                                                                                                                                  GGCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCCTTCTACTGCCAA
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                                                                   CCATGCCTAGACTGCGGGGCCCTGCACCGCCACACGCCTACTCTGTTCCCGCAGAGAT
                                                                                                                                   GGCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGCCAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Insert Length: 1568 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tumor Gene Index Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 698)
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National Cancer Institute, Cancer Genome Anatomy
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/db_xref="taxon:9606"
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/lab_host="DH10B"
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2.3e-134;
nes 7;
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CTTGCACGGAGCCCTGCGGCAACTCCACCTGCCTTGTGTGTCCCCAAGACACCTTCTTGG 181
                                                                            CAAGAAGATTGGTCTGTTTTGTTGCAGAGGCTGCCCAGC-GGGGCACTACCTGAAGGCCC 121
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                                                              CAAGAAGATTGGTCTGTTTGTTGCAGAGGCTGCCCAGCTGGGGGCACTACCTGAAGGCCC
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                                                                                                                                                                                     549;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 767)
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603065172F1 NIH_MGC_118 Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        High quality sequence stop: Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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                                                                                                                                                                                                                                                                                                                                    /note="Vector: pCMV-SPORT6; Site_1: NotI; Site_2: ECORV (destroyed); RNA source leukocytes from anonymous pool of non-activated adult donors. Library is oligo-dT primed and directionally cloned (EcoRV site is destroyed upon cloning). Average insert size 1.7 kb, insert size range
                                                                                                                                                                                                                                                            1.2-3.3 kb. Library is normalized and enriched for full-length clones and was constructed by C. Gruber (Invitrogen). Research Genetics tracking code 027. Note: this is a NIH_MGC Library."

243 c 241 g 161 t
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/clone_lib="NIH_MGC_118"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/db_xref="taxon:9606"
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99.1%;
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Pred. No. 3.1
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3.8e-132;
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                                                                                                                                                                                                                   Insert Length: 2028 Std Error:
Seq primer: -40UP from Gibco
High quality sequence stop: 445.
                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov Tissue Procurement: Ash Alizadeh,
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tg19b08.x1 NCI_CGAP_CLL1
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2.:, mRNA sequence.
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CDNA Library Preparation: M. Bento Soares, Ph.D.
CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome
Clone distribution: NCI-CGAP clone distribution
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tumor Gene Index
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/Clone_lib="NCI_CGAP_CLL1"
//tissue_type="B-cell, chronic lymphotic leukemia"
//lab_host="DH10B"
//note="Vector: pT7T3D-Pac (Pharmacia) with a modified
polylinker; Site_1: Not I; Site_2: Eco RI; 1st strand
was primed with a Not I - oligo(dT) primer [5]
                                                                                                                      /clone="IMAGE:2109207"
                                                                                                                                            /db_xref="taxon:9606"
                                                                                                                                                                                 1. .688
                                                                                                                                                             organism="Homo sapiens"
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Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
This clone is available royalty-f
                                                                                     Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eukaryota; Metazoa; Primates; Catarrhini; Hominidae; 1 (bases 1 to 562)
                                                           NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
National Cancer Institute, Cancer Genome Anai
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                        CCCACGAGCACCCTGGGGAGCTGTCCAGAGCGC
                                                                                                  ACTGACTGTGGGACCTGCCTGCCTTGCCTTCTATGAACATGGCGATGGCTGCGTGTCCTGC 480
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                    GCCTGGGAGAACCACATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGCC
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Plate:
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Tel: +82-42-860-4470
Fax: +82-42-860-4409
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Korea Research Institute of Bioscience & Biotechnology
52 Eoeun-dong Yuseong-gu, Daejeon 305-333, South Korea
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Kim, N.S., Hahn, Y.,
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21C Frontier Korean EST Project 2001
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                                                                                                                                                                                                                                                                                                                                                                                                                        /note="Organ: Stomach; Vector: pcNS; Site_1: EcoRI; Site_2: NotI; The poly (A)+ RNA was dephosphorylated with bacterial alkaline phosphatase (BAP) and then decapped with tabacco acid pyrophosphatase (TAP). The decapped intact mRNA was ligated with DNA-RNA linker including EcoR I site by treatment of T4 RNA ligase and the first strand cDNA was synthesized from ollog dT-selected mRNA by priming with dT-tailed vector. The dT-tailed vector was adjusted to have about 60nt. The cDNA vector was circularized with E. coli DNA ligase after digestion of EcoRI which site is also included in vector. An RNA strand competed to a DNA strand by Okayama-Berg method. The obtained cDNA vectors were used for transformation of competent cells E. coli TOPOIDF by electroporation method. The cDNA libraries constructed by this method are full-length enriched cDNA library."
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31 row: H column: 06
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/cell_line="SNU-620"
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3 similar to TR:000276 C00276 LYMPHOCYTE ASSOCIATED RECEPTOR OF DEATH 2. ; mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                        This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Possible reversed clone: similarity on wrong strand Possible reversed clone: polyT not found Seq primer: -40UP from Gibco High quality sequence stop: 489.
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Plate:
                                                                                                                                                                                      52 Eoeun-dong Yuseong-gu,
Tel: +82-42-860-4470
Fax: +82-42-860-4409
                                                                                                                                                                                                             Korea Research Institute of Bioscience & Biotechnology 52 Eoeun-dong Yuseong-gu, Daejeon 305-333, South Korea
                                                                                                                                                                                                                                                                                               Bukaryota; Metazoa; Chordata; Cranlata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 514)
Kim,N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim,M.R., Oh,K.J., Cheong,J.E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,S. and
                                                                                                                                                                                                                                                                                                                                                                                                            mRNA sequence.
BM756372
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27 row: G column: 03
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          /cell_type="Scattering floating"
/cell_line="SNU-620"
/lab_host="Top10F'"
                                                                                    /organism="Homo sapiens"
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/clone="S6SNU620-27-G03"
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K-EST0062030 S6SNU620 Homo sapiens

514 bp

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S6SNU620-32-A05

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Homo ....
Eukaryota; Metazur.
Mammalia; Eutheria; P
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N.S., Hahn,Y., C

Oh, J.H.,

Lee, J.Y.,

Ahn, H.Y.,

Chu, M.Y.,

Kim, M.R.

Metazoa; Chordata; Craniata; Vertebrata; Eutheria; Primates; Catarrhini; Hominidae;

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Pred. No. 5.1e-119;
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Park, H.S., Kim, S.

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21C Frontier Korean EST Project 2001
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Plate: 32 row: A column: 05
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L: +82-42-860-4470
c: +82-42-860-4409
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Site_2: NotI; The poly (A)+ RNA was dephosphorylated with
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K-EST0017828 S6SNU620 Homo sapiens
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Plate: 17 row: C column: 06
High quality sequence stop: 565.
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Fax: +82-42-860-4409
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                       Similarity
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    Conservative
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                                                                                             /note="Organ: Stomach; Vector: pCNS; Site_1: EcoRI; Site_2: NotI; The poly (A)+ RNA was dephosphorylated with bacterial alkaline phosphatase (BAP) and then decapped with tabacco acid pyrophosphatase (TAP). The decapped intact mRNA was ligated with DNA-RNA linker including EcoR I site by treatment of T4 RNA ligase and the first strand cDNA was synthesized from oligo dT selected mRNA by priming with dT-tailed vector. The dT-tailed vector was adjusted to have about 60nt. The cDNA vector was circularized with E. coli DNA ligase after digestion of EcoRI which site is also included in vector. An RNA strand converted to a DNA strand by Okayama-Berg method. The obtained cDNA vectors were used for transformation of competent cells E. coli ToplOF' by electroporation method. The cDNA libraries constructed by this method are full-length enriched cDNA library.*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone_lib="S6SNU620"
/sex="F"
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/cell_line="SNU-620"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           'lab_host="Top10F'"
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                     87.48;
98.68;
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Score 484.8; DB 14;
Pred. No. 1.1e-118;
0; Mismatches 7;
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                                                                                                                                                                                                        Unpublished (1997)
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                                                                                                                                                                                     Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                Contact: Robert Strausberg,
                                                                                                                                                                                                                               NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                 This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Insert Length: 1699 Std Error: 0.00
                                                                                                                                          primer:
                                                                                                                     quality sequence stop: 442.
Location/Qualifiers
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/organism="Homo sapiens"
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              Korea Research Institute | 52 Eoeun-dong Yuseong-gu, Tel: +82-42-860-4470 Fax: +82-42-860-4409
                                                                              Unpublished (2002)
                                                                                                                         Kim, N.S.,
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                                                           Genome Research Center
                                                                     Contact: Kim YS
                                                                                                             Oh, K.J., Cheong, J.E.,
                                                                                                                                                        Eukaryota; Metazoa;
                                                                                                                                                                   Homo sapiens
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(bases 1 to 492) m, N.S., Hahn, Y.,

Oh, J.H.,

,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y.,
Sohn,H.Y., Kim,J.M., Park,H.S., Kim,

Eutheria;

Chordata; Craniata; Vertebrata; Primates; Catarrhini; Hominidae;

Hominidae;

Homo

Kim,

Euteleostomi;

GI:19062345

S6SNU620 Homo

492 bp sapiens

mRNA cDNA clone

linear

S6SNU620-2-C03

EST 01-MAR-2002

Frontier Korean

EST Project 2001

of Bioscience & Biotechnology , Daejeon 305–333, South Korea

yongsung@mail.kribb.re

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CACAAGAAGATTGGTCTGTTTTGTTGCAGAGGCTGCCCAGCGGGGCACTACCTGAAGGCC
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ACTGACTGTGGGACCTGCCTGCCTGCCTTCTATGAACATGGCGATGGCTGCGTGTCC
                  CCATGCCTAGACTGCGGGGCCCTGCACCGCCACACGCCTACTCTGTTCCCGCAGAGAT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          85.9%;
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Plate: 21 row: D column: 12
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Tel: +82-42-860-4470
Fax: +82-42-860-4409
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                  remaining DNA into competent cells E. electroporation method."
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Query Match Best Local Similarity Matches 466; Conserv

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AGENCOURT_6794093 NIH_MGC_121
5', mRNA sequence.
BQ068309 GI:19897355
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Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information cal
                                                                                                                                                                                                                                                                                    Contact: Robert Strausberg, Ph.D.
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National Institutes of Health, Mammalian
Unpublished (1999)
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                                                                                                                                         found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
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                                                              /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5770562"
/clone_1ib="NIH_MGC_121"
/lab_host="DH10B"
/note-*Organ: brain; Vector: pCMV-SPORT6; Site_1: NotI; Site_2: EcoRV (destroyed); RNA source anonymous pool of fetal brains, female age 20 weeks, female age 24 weeks, and male age 26 weeks. Library is oligo-dT primed and directionally cloned (EcoRV site is destroyed upon
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48 CAGGGCACTACCTGAAGGCCCCTTGCACGGAGCCCTGCGGCAACTCCACCTGCCTTGTGT 107
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AW002222
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478 bp mRNA linear ES'
wt85c09.xl Soares_thymus_NHFTh Homo sapiens cDNA clone
IMAGE:2514256 3' similar to TR:014866 014866 SOLUBLE DI
                                              Eukaryota: Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 478)
                                                                                                                                      Tumor Gene Index Unpublished (1997)
                                                                                                                                                                                                                                                                                                     AW002222.1
EST.
                                                                                                     Contact: Robert Strausberg, I
Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                          NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
                                                                                                                                                                                                                                                                  Homo sapiens
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Location/Qualifiers
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/organism="Homo sapiens"

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Search completed: April Job time : 1742 secs
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Best Local Similarity 99.3
Matches 446; Conservative
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/clone_lib="Soares_thymus_NHFTh"
/dev_stage="fetal"
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Database
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Maximum Match 100%
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Perfect score:
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555
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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## SUMMARIES

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9 171	350.6	7 350.6	350.6	350.6	350.6	486.4	2 486.4	1 486.4	3 486.4	9 486.4	3 486.4	7 541	555	555	555	3 555	2 555	1 555	Score
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10797	433	433	433	433	433	1438	1438	1438	1438	1438	816	1743	1783	1634	1634	1634	1634	1254	Query Match Length
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US-10-092-154-1577	US-09-993-234-2	US-09-884-733-2	US-10-112-193-2	US-10-112-793-2	US-10-081-280-2	US-09-993-234-5	US-09-884-733-5	US-10-112-193-5	US-10-112-793-5	US-10-081-280-5	US-09-964-824A-292	US-09-954-531-1389	US-09-333-966-1	US-09-993-234-9	US-10-112-193-11	US-10-112-793-9	US-10-081-280-9	US-09-333-966-3	ID
Sequence 1577, Ap			Sequence 2, Appli		Sequence 2, Appli	Sequence 5, Appli	Sequence 5, Appli	Sequence 5, Appli .	Sequence 5, Appli	Sequence 5, Appli	Sequence 292, App	œ	•	•	_		Sequence 9, Appli	Sequence 3, Appli	Description

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31.6	31.6	31.6	31.6	32	33.2	33.2	33.2	. 34.4	34.6	34.8	34.8	34.8	34.8	34.8	34.8	34.8	35	36.4	37.4	38	40.4	42.2	42.2	55.4	171
5.7	5.7	5.7	5.7	5.8	6.0	6.0	_	6.2				6.3	6.3	6.3	6.3	6.3	6.3	6.6	6.7	6.8	7.3	7.6	7.6	10.0	30.8
1801	746	518	299	1780	155074	37113	15540	896	20556	5988	4898	4898	2173	2173	2173	2173	1345	301	5870	2130	671	2440	543	346	10797
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US-09-822-830A-128	US-09-910-943-165	US-09-864-761-8725	US-09-864-761-25435	US-09-917-800A-1521	US-10-026-188-6	US-09-999-121-11	US-09-999-121-12	US-09-884-441-22	US-09-880-107-3945	US-10-044-090-90	US-09-880-107-2323	US-09-919-497-35	US-09-899-422-14	US-09-792-356-14	US-09-899-429A-24	US-09-898-234-14	US-09-853-450-7	US-10-070-676-7	US-09-838-718A-8	US-09-917-800A-1601	US-10-184-644-346	US-09-970-532-1	US-09-970-532-3	US-09-783-590-1252	US-09-764-847-1577
Sequence 128, App		Sequence 8725, Ap	Sequence 25435, A	Sequence 1521, Ap	Sequence 6, Appli	Sequence 11, Appl	Sequence 12, Appl			Sequence 90, Appl	Sequence 2323, Ap	Sequence 35, Appl	9 14	Sequence 14, Appl	Sequence 24, Appl	14,	Sequence 7, Appli	Sequence 7, Appli	Sequence 8, Appli	Sequence 1601, Ap	Sequence 346, App	Sequence 1, Appli	Sequence 3, Appli	Sequence 1252, Ap	Sequence 1577, Ap

## ALIGNMENTS

RESULT 1

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US-09-333-966-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 3, Application US/09333966 Patent No. US20020009773A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/333,966
FILING DATE:
OTRECTION DATE:
                                                                                           APPLICATION NUMBER: US/08/815,469
FILING DATE:
APPLICATION NUMBER: No. US20020009773A1 Yet Assigned FILING DATE: 06-FEB-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/028,711
FILING DATE: 17-OCT-1996
PRIOR APPLICATION DATA:
FILING DATE: 17-OCT-1996
PRIOR APPLICATION DATA:
FILING DATE: 17-OCT-1996
PRIOR APPLICATION DATA:
                       APPLICATION NUMBER: US 60/013,285
FILING DATE: 12-MAR-1996
ATTORNEY_AGENT INFORMATION:
NAME: Steffe, Eric K.
                                                                                                                                                                                                                                                                                            CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Dixit, Vishva
APPLICANT: Gentz, Reiner L.
APPLICANT: Dillon, Patrick J.
TITLE OF INVENTION: Death Domain
NUMBER OF SEQUENCES: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Yu, Guo-
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
ZIP: 20005-3934
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Washington
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                     Steffe,
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1100 New York Ave.,
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36,688
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Sequence 9, Application US/10081280
Patent NO. US20020165157A1
GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-2 LI AND AP.
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS: Genentech, Inc.
STREET: 460 Point San Bruno B1
CITY: South San Francisco
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US-10-081-280-9
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Best Local :
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INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
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TELECOMMUNICATION INFORMATION: 202-371-2600
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STRANDEDNESS: double
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Pred. No. 3e-161;
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Best Local S
Matches 555
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TELEPHONE: 415/225-5416
TELEPHONE: 415/25-9881
TELEFAX: 415/952-9881
TELETAX: 910/371-7168
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 1634 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                             626
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APPLICATION NUMBER: US/10/081,280
FILING DATE: 21-Feb-2002
                                                                                                                           CCATGCCTAGACTGCGGGGCCCTGCACCGCCACACACGGCTACTCTGTTCCCGCAGAGAT
                                                                                                                                                                        GGCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCCTTCTACTGCCAA
                                CCATGCCTAGACTGCGGGGCCCTGCACCGCCACACGGCTACTCTGTTCCCGCAGAGAT
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MEDIUM TYPE: 3.5 i
                                                                                                                                                            GGCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGCCAA
                                                                                                                                                                                                         TCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGAAGCCA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: PI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 21-Feb-2002 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/0 FILING DATE: 31-Mar-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION DATA:
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Pred. No. 3.2e-161;
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                                                                                                                                                                                                                                                                                                                                              Local
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TELEPHONE: 650/225-5416
TELEFAX: 650/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Marschang, Diane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 94080
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 mb floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Widpatin (Genentech)
CURRENT APPLICATION DATA:
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ADDRESSEE: Genentech, Inc.
STREET: 1 DNA Way
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Ashkenazi, Avi J.
                                 CCTTGCACGGAGCCCTGCGGCAACTCCACCTGCCTTGTGTGTCCCCAAGACACCTTCTTG 180
                                                                                                                                                                             CACAAGAAGATTGGTCTGTTTTGTTGCAGAGGCTGCCCAGCGGGGCACTACCTGAAGGCC
                                                                                                                                                                                                  CACAAGAAGATTGGTCTGTTTTGTTGCAGAGGCTGCCCAGCGGGGCACTACCTGAAGGCC 120
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TCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCCA
                                                                     GCCTGGGAGAACCACCATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGCC 240
                                                                                                                                                                                                                                                                                                                            555;
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FILING DATE: 28-Mar-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Marschang, Diane L. REGISTRATION NUMBER: 35,600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 08/625328
FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/828,683A FILING DATE: 31-Mar-1997
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Pred. No. 3.2e-161;
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; SEQUENCE DESCRIPTION: SEQ ID NO: US-10-112-193-11
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Publication No. US20030004313A1
GENERAL INFORMATION:
                      Query Match
Best Local :
     Matches
                                                                                                                                                  INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 1634 base pairs
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   Local Similarity
les 555; Conserv
                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: p1052R1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Apo-3 POLYPEPTIDE NUMBER OF SEQUENCES: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Ashkenazi, Avi J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ACTGACTGTGGGACCTGCCTGCCTGCCTTCTATGAACATGGCGATGGCTGCGTGTCCTGC
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FILING DATE: 28-Mar-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech)
                                                                                                                  TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                          NAME: Marschang, Dia
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 08/928,069 FILING DATE: 11-Sep-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: South San F
STATE: California
                                                                                                                                                                                                        TELEFAX: 650/952-9881
                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 60/026943 FILING DATE: 23-Sep-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET:
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 100.0%; ilarity 100.0%; iconservative 0;
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MBER: 35,600
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 Score 555; DB 9;
Pred. No. 3.2e-161;
; Mismatches 0;
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Gaps
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Length 1634;

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RESULT 5
US-09-993-234-9
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Patent No. US20020146768A1
GENERAL INFORMATION:
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                NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-5416
                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/828,683
FILING DATE: <UNKNOWN>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 i
                                                                                                                                                                                       CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-2 LI AND
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                                                                                                                                                                                                                                                                                                    STREET: 460 Point San Bruno Blvd CITY: South San Francisco STATE: California
                                                                                                                                                     APPLICATION NUMBER: US/09/993,234 FILING DATE: 19-NO. US20020146768A1-2001
                                                                                                                                                CLASSIFICATION:
                                                                                                                                                                                                                                                                                     COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Genentech, Inc.
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                                                                                                                                                                                           Sequence 1, Application US/09333966 Patent No. US20020009773A1
                                                                                       APPLICANT: Yu, Guo-Liang
APPLICANT: Ni, Jian
APPLICANT: Dixit, Vishva
APPLICANT: Gentz, Reiner L.
APPLICANT: Dillon, Patrick J.
TITLE OF INVENTION: Death Domai
                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Yu, Gu
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Best Local 9
STREET: 1100
CITY: Washington
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                                                                 CORRESPONDENCE ADDRESS:
                                                                                NUMBER OF SEQUENCES:
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                                                     ADDRESSEE:
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TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
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TELEX: 910/371-7168
                                     E: Sterne, Kessler,
1100 New York Ave.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; ilarity 100.0%; Conservative
                                                                              Death Domain Containing
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Pred. No. 3.2e-161;
; Mismatches 0;
                                       Goldstein
NW, Suite
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COUNTRY: ZIP: 200

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US-09-333-966-1
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Best Local Sim
Matches 555;
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APPLICATION NUMBER: NO. US200200097
FILING DATE: 06-FEB-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/028,711
FILING DATE: 17-OCT-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/013,285
PRIOR APPLICATION NUMBER: US 60/013,285
APPLICATION NUMBER: 36,688
APPLICATION NUMBER: 36,688
APPLICATION NUMBER: 36,688
APPLICATION NUMBER: 36,688
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INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 141
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
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hes 555; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid STRANDEDNESS: double TOPOLOGY: both
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CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                 CTGGGGGCCCGGGCCCAGGGCGCACTCGTAGCCCCAGGTGTGACTGTGCCGGTGACTTC 60
CCATGCCTAGACTGCGGGGCCCTGCACCGCCACACACGGCTACTCTGTTCCCGCAGAGAT
                                                                      GGCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGCCAA 360
                                                                                                                                TCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGGCTGTAAGCCA
                                                                                                                                                                         GCCTGGGAGAACCACCATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGCC
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                                                                                                                 TCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCCA
                                                       GGCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGCCAA
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Pred. No. 3.2e-161;
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CURRENT FILING DATE: 2002-05-02
PRIOR APPLICATION NUMBER: US/60/233,133
PRIOR FILING DATE: 2000-09-18
PRIOR APPLICATION NUMBER: US/60/234,009
PRIOR APPLICATION NUMBER: US/60/234,034
PRIOR APPLICATION NUMBER: US/60/234,034
PRIOR APPLICATION NUMBER: US/60/234,034
PRIOR APPLICATION NUMBER: US/60/234,509
PRIOR APPLICATION NUMBER: US/60/234,509
PRIOR FILING DATE: 2000-09-22
PRIOR FILING DATE: 2000-09-22
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SOFTWARE: Patentin version
SEQ ID NO 1389
LENGTH: 1743
TYPE: DNA
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APPLICANT: Weaver,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Process for Identifying Anti-Cancer Therapeutic Agents Using TITLE OF INVENTION: Gene Sets FILE REFERENCE: 689290-77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US/60/234,567
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CCATGCCTAGACTGCGGGGCCCTGCACCGCCACACACGGCTACTCTGTTCCCGCAGAGAT
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                                                                   GGCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCCTTCTACTGCCAA
                                                                                                           TCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGAAGCCA
                                                                                                                        TCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCCA
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Pred. No. 6.5e-157;
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SEQ ID NO 292
LENGTH: 816
TYPE: DNA
ORGANISM: Homo sapiens
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR FILING DATE: 2000-09-28
NUMBER OF SEQ ID NOS: 583
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PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR FILING DATE: 2000-09-28
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ACTGACTGTGGGACCTGCCTGCCTGCCTTGTGTATGAACATGGCGATGGCTGCGTGTCCTGC
                                   CCATGCCTAGACTGCGGGGCCCTGCACCGCCACACGGCTACTCTGTTCCCGCAGAGAT
                                               CCATGCCTAGACTGCGGGGCCCTGCACCGCCACACACGGCTACTCTGTTCCCGCAGAGAT
                                                                                                                                              TCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGGCTGTAAGCCA
                                                                                                                                                                                             GCCTGGGAGAACCACATAÀTTCTGAATGTGCCGGCTGCCAGGCCTGTGATGAGCAGGCC
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98.8%;
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Pred. No. 3.6e-140;
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TOPOLOGY: Linear; SEQUENCE DESCRIPTION: US-10-081-280-5
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US-10-081-280-5
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                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                 434 CTGGGGGCCCGGGCCACGGCACTCGTAGCCCCAGGTGTGACTGTGCCGGTGACTTC 493
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TCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCCGCTGTGGCCTGTAAGCCA
                                                                                                                                 CCTTGCACGGAGCCCTGCGGCAACTCCACCTGCCTTGTGTGTCCCCCAAGACACCTTCTTG
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APPLICATION NUMBER: US/10/081,280
FILING DATE: 21-Feb-2002
CLASSIFICATION: <Unknown>
                                                        REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
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TITLE OF INVENTION: Apo-2 LI AND Apo-3 POLYPEPTIDES
NUMBER OF SEQUENCES: 11
                                                                                                                                                                                                         CACAAGAAGATTGGTCTGTTTTGTTGCAGAGGCTGCCCAGCGGGCACTACCTGAAGGCC 553
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TYPE: Nucleic Acid
STRANDEDNESS: Single
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Marschang, Diane L. REGISTRATION NUMBER: 35,600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/829,270 FILING DATE: 31-Mar-1997
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CITY: South San Francisco
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                                                                                                                                                                                                                                                                                                                                                                 Conservative
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98.8%;
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Pred. No. 3.
                                                                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                                                                           1; DB 9;
3.9e-140;
hes 6;
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TOPOLOGY: Linear SEQUENCE DESCRIPTION: SEQ ID NO: US-10-112-793-5
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US-10-112-793-5
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                                                                      Query Match
Best Local Similarity
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                                                           Matches
                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1438 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
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1 CTGGGGGCCCGGGCCCAGGGCGCACTCGTAGCCCCAGGTGTGACTGTGCCGGTGACTTC
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                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-2 LI AND Apo-3 POLYPEPTIDES
NUMBER OF SEQUENCES: 28
                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
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                                                          490;
                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/828,683A FILING DATE: 31-Mar-1997
APPLICATION NUMBER: 08/625328
FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
                                                                                                                                                        TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/10/112,793 FILING DATE: 28-Mar-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: 3.5 inch, 1.44 Mb: COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 1 DNA Way
CITY: South San Francisco
                                                                                                                                                                                                                                                                                        NAME: Marschang, Dian REGISTRATION NUMBER:
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                                                                                                                                                                                                                                 TELEFAX: 650/952-9881
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                                                          Conservative
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                                                                      87.6%;
98.8%;
                                                                                                                                                                                                                                                                                                    Diane L.
                                                       Score 486.4; DB 9;
Pred. No. 3.9e-140;
0; Mismatches 6;
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                                                                                  Length 1438;
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RESULT 11
US-10-112-193-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                   INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      914 CCCACGTAATTCCTAG
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FILING DATE: 23-Sep-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech) CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy
                                                       NAME: MAISCHANG, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1052R1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/928,069
FILING DATE: 11-Sep-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-3 POLYPEPTIDE
NUMBER OF SEQUENCES: 15
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/10/112,193 FILING DATE: 28-Mar-2002
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CITY: South
                                                                                                                                                                                                                                                                                          CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: California
                                       TELEFAX: 650/952-9881
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RESULT 12
US-09-884-733-5
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                                                                                                                                                                                                                                                                                                             Sequence 5, Application US/09884733
Patent No. US20020123116A1
GENERAL INFORMATION:
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Best Local (
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                                                                          MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech) CURRENT APPLICATION DATA:
                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 1
                     PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Apo-2 Ligand Inhibitor
                                                                                                                                                                                                                                                                                                APPLICANT: Ashkenazi, Avi J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CTGGGGGCCCGGGCCCAGGGCGGCACTCGTAGCCCCAGGTGTGACTGTGCCGGTGACTTC
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                              APPLICATION NUMBER: US/09/884,733
FILING DATE: 19-Jun-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                          ADDRESSEE: Genentech, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
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                                                                                                                                                                     94080
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98.8%;
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Pred. No. 3.9e-140;
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                                                                                                                            RESULT 13
US-09-993-234-5
; Sequence 5, Application US/09993234
; Patent No. US20020146768A1
; GENERAL INFORMATION:
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Best Local :
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          CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 460 Point San Bruno
CITY: South San Francisco
STATE: California
                                                                                     APPLICANT: Ashkenazi, Avi
TITLE OF INVENTION: Apo-2
NUMBER OF SEQUENCES: 11
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NAME: Marschang, Diane
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STATE: Ca
COUNTRY:
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TYPE: Nucleic Acid
STRANDEDNESS: Single
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TELEX: 91
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910/371-7168
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Pred. No. 3.
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Best Local Similarity
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 08/828,683 FILING DATE: <Unknown> ATTORNEY/AGENT INFORMATION:
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                                                                                                  GGCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGCCAA 360
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                                                             TCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCCA
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CCCACGTAATTCCTAG
                                                 ACTGACTGTGGGACCTGCCTGCCTTCTATGAACATGGCGATGGCTGCGTGTCCTGC
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STRANDEDNESS: Single
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FILING DATE: 19-No. US20020146768A1-2001
CLASSIFICATION: <UNKnown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007P1
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TELEFAX: 415/952-9881
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                                                                                                                                                                                                                                                                       LENGTH: 1438 base pairs
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98.8%;
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Pred. No. 3.9e-140;
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RESULT 14 US-10-081-280-2

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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
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FILING DATE: 31-Mar-1997
ATTORNEY_AGENT INFORMATION:
NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007R1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-5416
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CORRESPONDENCE ADDRESS:
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CAACCATGCCTAGACTGCGGGGCCCTGCACCGCCACACACGGCTACTCTGTTCCC 412
                                       PRIOR APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Ashkenazi, Avi
TITLE OF INVENTION: Apo-2
                                                                                                                                                                                                                                                                                      CACAAGAAGATTGGTCTGTTTGGTGCAGAGGCTGCCCAGCGGGGCAACTACCTGAAGGC 123
                                                                                                                                                                                                                                                                                                                                                   CTGGGGGCCCGGGCCAGNGGCGGCACTCGTAGCCCCAGGTGTGACTGTGCCGGTGACTTC 63
                                                                                                                                                                GGCCTGGGAGAACCACCATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGC 239
                                                                                                                                                                                                                           CCCTTGCACGGAGCCCTGC-GCAACTCCACCTGCCTTGTGTGTCCCCAAGACACCTTCTT
                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 433 base pairs TYPE: Nucleic Acid STRANDEDNESS: Single TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/10/081,280 FILING DATE: 21-Feb-2002 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech)
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CITY: South San Francisco
STATE: California
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96.48;
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RESULT 15 US-10-112-793-2

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TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
TELEPAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 433 harm
                                                                                                                                                                                                                                                                                                                                                                                 Query Match 63.2%;
Best Local Similarity 96.4%;
Matches 400; Conservative (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-2 LI AND Apo-3 POLYPEPTIDES
NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
   243
                                                                                              180 GGCCTGGGAGAACCACATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGC 239
                                                                                                                                                        124
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
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CCCTTGCACGGAGCCCTGC-GCAACTCCACCTGCCTTGTGTGTCCCCAAGACACCTTCTT
                                                                           GGCCTGGGAGAACCACCATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 433 base pairs
TYPE: Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/828,683A FILING DATE: 31-Mar-1997 APPLICATION NUMBER: 08/625328 FILING DATE: 1-Apr-1996 APPLICATION NUMBER: 08/710802 FILING DATE: 23-Sep-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/10/112,793
FILING DATE: 28-Mar-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Marschang, Diane L. REGISTRATION NUMBER: 35,6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 94080
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STREET: 1 DNA Way
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Search completed: April Job time : 128 secs 6, 2003, 16:30:18 밁

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Perfect score:
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Maximum Match
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Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Issued_Patents_NA:*
1: /cgn2_6/ptodata/1
2: /cgn2_6/ptodata/1
3: /cgn2_6/ptodata/1
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Match
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Listing first 45 summaries
   GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
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US-08-928-69-5
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US-08-928-683A-2
US-09-513-007-3
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US-09-513-007-1
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US-09-404-879A-23
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US-08-402-066-3
US-08-402-066-3
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GENERAL INFORMATION:
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                                                        TELEFAX: 202-371-2540 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Yu, Guo-
APPLICANT: Ni, Jian
APPLICANT: Dixit, V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                       CITY: Washington
STATE: DC
                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
            STRANDEDNESS:
                                                                                TELEPHONE:
                                  LENGTH:
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                                  1254 base pairs
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Result No.

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5.6	5.6	5.6	5.6	5.6	5.6	5.6	5.6	5.6	ა ნ	5. 6	5.6	5.6	5.6	5.6	5.9	6.0	6.0
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US-08-142-368A-8	US-08-560-024-1	US-08-299-849B-8	US-08-190-411A-1	US-07-807-043B-8	US-08-627-820-23	US-09-056-105-4	US-08-465-167A-23 .	US-08-037-230D-7	US-08-967-727-7	US-08-142-368A-7	US-08-299-849B-7	US-07-807-043B-7	US-08-845-528C-8	US-08-993-118-8	US-09-221-017B-687	US-09-282-357-8	US-08-918-288-8
Sequence 8, Appli	Sequence 1, Appli	Sequence 8, Appli	Sequence 1, Appli	Sequence 8, Appli	Sequence 23, Appl	Sequence 4, Appli	Sequence 23, Appl	Sequence 7, Appli	Sequence 8, Appli	Sequence 8, Appli	Sequence 687, App	Sequence 8, Appli	Sequence 8, Appli				

ALIGNMENTS

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/815,469
FILING DATE: HEREWITH

CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: No. 6153402 Yet
FILING DATE: 06-FEB-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/028,711
FILING DATE: 17-OCT-1996
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA: ZIP: 20005-3934 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk APPLICATION NUMBER: US 60/013,285
FILING DATE: 12-MAR-1996
ATTORNEY/ACENT INFORMATION:
NAME: Steffe, Eric K. SEQUENCE CHARACTERISTICS: TELECOMMUNICATION INFORMATION: APPLICANT: Dillon, Patrick J.
TITLE OF INVENTION: Death Domain Containing Receptors
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS: COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30 REGISTRATION NUMBER: 36 REFERENCE/DOCKET NUMBER: Application US/08815469 E: Sterne, Kessler, 1100 New York Ave., Dixit, Vishva Gentz, Reiner L. Yu, Guo-Liang 202-371-2600 36,688 **س** 6153402 Yet Assigned 1488.0310003/EKS/KRM Goldstein & Fox, P.L.L.C. NW, Suite 600

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; LOCATION:
US-08-815-469-3
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Best Local S
Matches 555
                                                                                                                                                              Patent No.
GENERAL II
    COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb f
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                    APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-3 POLYPEPTIDE
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
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                                                                    ADDRESSEE:
STREET: 1
CITY: Sout
STATE: Cal
COUNTRY: (
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APPLICATION
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                                                                                                                                                                        11, Application US/08928069
o. 6462176
                                                                                                                                                             INFORMATION:
                                                                 94080
                                                                                             South San Francisco
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                                                                                                             Genentech, Inc.
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US/08/928,069
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Pred. No. 8.5e-148;
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RESULT 3
US-08-828-683A-9
; Sequence 9, Application US/088286; Patent No. 6469144; GENERAL INFORMATION: APPLICANT: AShkenazi, Avi; TITLE OF INVENTION: Apo-2; NUMBER OF SEQUENCES: 28
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REGISTATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1
TELECOMMUNICATION INFORMATION:
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FILING DATE: 09/23/1996
ATTORNEY/AGENT INFORMATION:
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TELEFAX: 650/952-9881
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DEDNESS: Single
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N: 435
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Pred. No. 9.3e-148;
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US-08-828-683A-9
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Local Similarity 100.0%;
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NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
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LENGTH: 1634 base pairs
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
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                 ACTGACTGTGGGACCTGCCTGCCTGCCTTCTATGAACATGGCGATGGCTGCGTGCCTGC 480
                                                                                                         CCATGCCTAGACTGCGGGGCCCTGCACCGCCACACGCGCTACTCTGTTCCCGCAGAGAT 420
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                                                                                                                                           GGCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGCCAA
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ACTGACTGTGGGACCTGCCTGCCTGCCTTCTATGAACATGGCGATGGCTGCGTGTCCTGC
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FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
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Pred. No. 9.3e-148;
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US-08-815-469-1
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                                                         Query Match
Best Local Similarity
Matches 555; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/
FILING DATE: HEREWITH
CLASSIETCATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: NO.
                                                                                                                                                                                                                                                                                                                      ATTORNEY AGENT INFORMATION:
NAME: Steffe, Eric K.
REGISTRATION NUMBER: 36,0
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: 202-371-2600
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APPLICATION NUMBER: US 60/028,711
FILING DATE: 17-0CT-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Death Domain Containing Receptors NUMBER OF SEQUENCES: 17
                                                                                                                                                           MOLECULE TYPE: FEATURE:
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                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 60/013,285 FILING DATE: 12-MAR-1996
                                                                                                                                             NAME/KEY:
                                                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                  LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
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               1 CTGGGGGCCCGGGCCCAGGGCGCACTCGTAGCCCCAAGGTGTGACTGTGCCGGTGACTTC 60
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Gentz, Reiner L.
Dillon, Patrick
                                                         Conservative
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                                                      Score 555; DB 3;
Pred. No. 9.6e-148;
; Mismatches 0;
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NW, Suite
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                                                                                   Length 1783;
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US-08-928-069-5
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                    TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb f
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
                                             REFERENCE/DOCKET NUMBER: P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
                                                                                                                 APPLICATION NUMBER: 60/02 FILING DATE: 09/23/1996 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                          CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Ashkenazi, Avi
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CORRESPONDENCE ADDRESS
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                                                                                   NAME: Marschang, Diane L. REGISTRATION NUMBER: 35,600
                                                                                                                                                                                                                                                                                                                                STREET: 1 DNA WE
CITY: South San
STATE: Californi
                                                                                                                                                                                   APPLICATION NUMBER: US/08/928,069
FILING DATE: 11-Sep-1997
                                                                                                                                                                                                                                                                                                      COUNTRY: UZIP: 94080
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        CHARACTERISTICS
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1 DNA Way

San Francisco
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                                                                                                                                                                                                                                                                                                                                California
                                 650/952-9881
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US-08-828-683A-5
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// STRANDEDNESS: Sit
// TOPOLOGY: Linear
US-08-928-069-5
                                                                                                                                                                                                                                                 Sequence 5, Application US/08828683A
Patent No. 6469144
GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-2 LI AND
NUMBER OF SEQUENCES: 28
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Best Local Similarity
                                 COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION NUMBER: US/08/828,683A
FILING DATE: 31-Mar-1997
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           CLASSIFICATION: <Unknown> PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
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                                                                                                                                                                 ADDRESSEE: Genentech, 1
STREET: 1 DNA Way
CITY: South San Francis
STATE: California
COUNTEX: USA
                                                                                                                                                         ZIP: 94080
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98.8%;
NUMBER:
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Apo-3 POLYPEPTIDES

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                                   CACAAGAAGATTGGTCTGTTTTGCTGCAGAGGCTGCCCAGCGGGGCACTACCTGAAGGCC 120
                                                                                                              Score 486.4;
Pred. No. 2.
                                                                                                    Mismatches
                                                                                                     ..2e-128;
ies 6;
                                                                                                                        DB 4;
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                                                                                                      Indels
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FILING DATE: 1-Apr-1996 APPLICATION NUMBER: 08/710802 FILING DATE: 23-Sep-1996

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Best Local Sim
Matches 490;
                                                                                         Sequence 2, Application US/08928069
Patent No. 6462176
GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-3 POLYP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1438 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
STREET: 1 L...
CITY: South San Fre
                                                           NUMBER OF INVENTION: APO-3
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
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                                                    ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION: NAME: Marschang, Diane
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                                                   Genentech, Inc.
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                                                                                               POLYPEPTIDE
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Pred. No. 2.2e
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RESULT 8 US-08-,828-683A-2

Sequence 2, Application US/08828683A
Patent NO. 6469144
GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-2 LI AND Apo-3 POLYPEPTIDES
NUMBER OF SEQUENCES: 28

CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech,
STREET: 1 DNA Way

Inc.

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Best Local Similarity 96.4
Matches 400; Conservative
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NAME: MAISCHANG, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P105
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 11-Sep-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/0
FILING DATE: 09/23/1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech) CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 inch, 1.44 mb floppy disk
 363
                                                                        300 AGGCTGGTTTGTGGAGTGCCA-GGTCAGCCAATGTGTCAGCAG-TTCACCCCTTCTACTGC
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STRANDEDNESS: Single
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4 CTGGGGGCCCGGGCCAGNGGCGGCACTCGTAGCCCCAGGTGTACTGTGCCGGTGACTTC
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GGCCTGGGAGAACCACCATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGC
                               CAACCATGCCTAGACTGCGGGGCCCTGCACCGCCACACACGGCTACTCTGTTCCC 412
                                                           GGGCTGGTTTGTGGAGTGCCAGGGTCAGCCAATGTGTCAGCAGTTTCACCCCTTCTAATGC
                                                                                                                    CTCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGGTTGGCTGTAAGCA
                                                                                                                                                                                GGCCTGGGAGAACCACCATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGC
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ER: P1052R1
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 350.6;
Pred. No. 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                          6; DB 4;
3.6e-90;
hes 11;
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COMPUTER READABLE FORM:

CITY: South San F STATE: California COUNTRY: USA

Francisco

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US-09-513-007-3
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               Sequence 3, Application US/09513007 Patent No. 6406907 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches 400;
APPLICANT: Taylor, J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                             363 CAACCATGCCTAGACTGCGGGGCCCTGCAACGCAACACACGGCTAATNTGTTTCC 417
                                                                                                                                                                    358
                                                                                                                                                                                                                    300 AGGCTGGTTTGTGGAGTGCCA-GGTCAGCCAATGTGTCAGCAG-TTCACCCTTCTACTGC
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APPLICATION UNMBER: 08/625328

FILING DATE: 1-Apr-1996

APPLICATION NUMBER: 08/710802

FILING DATE: 23-Sep-1996

ATTORNEY/AGENT INFORMATION:

NAME: MAISCHAMP, Diane L.

REGISTRATION NUMBER: 35,600

REFERENCE/DOCKET NUMBER: P1007E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 CTGGGGGCCCGGGCCCAGGGCGCACTCGTAGCCCCAGGTGTGACTGTGCCGGTGACTTC 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/828,683A
FILING DATE: 31-Mar-1997
CLASSIFICATION: <Unknown>
                                                                                                                                                    CAACCATGCCTAGACTGCGGGGCCCTGCACCGCCACACACGGCTACTCTGTTCCC 412
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
TELEFAX: 650/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE DESCRIPTION: SEQ ID
                                                                                                                                                                                                GGGCTGGTTTGTGGAGTGCCAGGGTCAGCCAATGTGTCAGCAGTTTCACCCTTCTAATGC
                                                                                                                                                                                                                                                                                                                                  GGCCTGGGAGAACCACCATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                     CCCTTGCACGGAGCCCTGCGGCAACTCCACCTGCCTTGTGTGTCCCCCAAGACACCTTCTT 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CACAAGAAGATTGGTCTGTTTTGCAGAGGCTGCCCAGCGGGGCAACTACCTGAAGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CACAAGAAGATTGGTCTGTTTTGTTGCAGAGGCTGCCCAGCGGGGC-ACTACCTGAAGGC 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTGGGGGCCCGGGCCAGNGGCGGCACTCGTAGCCCCAGGTGTGACTGTGCCGGTGACTTC 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             63.2%;
illarity 96.4%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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Pred. No. 3.6e-90;
D; Mismatches 11;
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GENERAL INFORMATION:

APPLICANT: Taylor, J. Michael

APPLICANT: Kehrli, Jr., Marcus

APPLICANT: Lee, Eun-Kyung

APPLICANT: Lee, Eun-Kyung

APPLICANT: Mwangi, Simon

TITLE OF INVENTION: BOVINE TUMOR NECROSIS FACTOR RECEPTOR-1

TITLE OF INVENTION: AND METHODS OF USE

FILE REFERENCE: 08411-018001

CURRENT APPLICATION NUMBER: US/09/513,007

CURRENT APPLICATION NUMBER: 60/122,156

PRIOR APPLICATION NUMBER: 60/122,156

PRIOR APPLICATION NUMBER: 60/122,156

PRIOR FILING DATE: 1999-02-26
; NAME/KEY: CDS
; LOCATION: (294)...(1706)
US-09-513-007-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 10
US-09-513-007-1
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                                                                                                                                                              NUMBER OF SEQ ID NOS: 8
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 1, Application US/09513007 Patent No. 6406907
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 3
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TITLE OF INVENTION: BOVINE TUMOR NECROSIS FACTOR RECEPTOR-1
TITLE OF INVENTION: AND METHODS OF USE
FILE REFERENCE: 08411-018001
CURRENT APPLICATION NUMBER: US/09/513,007
CURRENT FILING DATE: 2000-02-25
CURRENT FILING DATE: 2000-03 166
                                                                                          LENGTH: 2440
TYPE: DNA
ORGANISM: Bos taurus
                                                                              FEATURE:
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PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 8
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TYPE: DNA
ORGANISM: Bos taurus
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LOCATION: (1)
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Local Similarity 50.2%;
les 132; Conserva+:.-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CCACCCGCAAAATAGCACCATTTGCTGCACCAAGTGCCACAAAGGTACCTATCTGTACAA 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   -TTCCAGGTGGAGATTTCGCCTTGTGTAGTGGACCGGGACACTGTGTGCGGCTGCAGGAA 299
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Pred. No. 0.0058;
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US-08-592-874-1/c
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Best Local Similarity
                                                                                               TELEFAX: 212-818-9479
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
TOPOLOGY: unknown MOLECULE TYPE: DNA (FRAGMENT TYPE: N-ter
                                                                                                                                             TELEPHONE: 212-986-4090
                                                                                                                                                                                                                             CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UDMBER: US 0:
FILING DATE: 24-JAN-1995
                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: DNA SEGMENTS AND METHODS FOR INCREASING TITLE OF INVENTION: POLYSACCHARIDE PRODUCTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          680 GAACCAGTACCGGGAATACTGGG 702
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   180
                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                 STRANDEDNESS:
                                                                                                                                                                               NAME: GOLDBERG, JULES E. REGISTRATION NUMBER: 24,
                                                                                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              60 CCACAAGAAGATTGGTCTGTTTTGCTTGCAGAGGCTGCCCAGCGGGGCACTACCTGAAGGC 119
                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
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                                                                                  LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TGCCTTGGAGAACCATCTCAGACGATGCCTGAGCTGCTCCAGGTGCCGGGACGAAATG---
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    Application US/08592874
    5854034

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                                                         nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   261 MADISON AVENUE
                                                                                                                                                                                                                                                                                                                                                                                                                                      USA
                                                                                                                                212-818-9479
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MIKOLAJCZAK, MARCIA
ARMENTROUT, RICHARD W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                THORNE, LINDA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 YAMAZAKI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                POLLOCK, THOMAS J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   JULES E.
      N-terminal
                  DNA (genomic)
                                                   unknown
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50.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GOLDBERG
                                                                                                                                                                                                                                                                                                         US/08/592,874
                                                                                                                                                                               24,408
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Pred. No. 0.0095;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               128;
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; ORGANISM: Sphingomonas
US-09-096-942-2
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Patent No. 6027925
GENERAL INFORMATION:
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                                SOFTWARE:
SEQ ID NO 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Production of Xanthan Gum by Sphingomonas Bacteria TITLE OF INVENTION: Carrying Genes from Xanthomonas Campestris FILE REFERENCE: seg list for appl filed from pro. appl CURRENT APPLICATION NUMBER: US/09/096,942

CURRENT FILING DATE: 1998-06-12
                                                                                                                            CURRENT FILING DATE: 1998-06-11
                                                                                                                                             TITLE OF INVENTION: Production of Xanthan Gum by Sphingomonas Bacteria TITLE OF INVENTION: Carrying Genes from Xanthomonas Campestris FILE REFERENCE: seel list for appl filed from pro. appl CURRENT APPLICATION NUMBER: US/09/096,867
                                                                                                                                                                                                                   APPLICANT: Thorne, Linda APPLICANT: Armentrout, Richard W
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APPLICANT: Thorne, Linda
APPLICANT: Armentrout, Richard W
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APPLICANT: Mikolajczak, Marcia
                                                                                                          EARLIER APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn
                                                                       NUMBER OF SEQ ID NOS: 2
                                                                                       EARLIER FILING DATE:
                                                                                                                                                                                                                                                                                                APPLICANT: Pollock, Thomas J
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TYPE: DNA
                   LENGTH:
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US-08-762-308-10
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Best Local Similarity
Matches 120; Conserv
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                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 1956 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 05-APR-1994 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7480 GTGCCGGCTGGGAGGTCTTCATGAAGAAATTGTACCTGGTAACGGCGGTGGCTGCGGCCG
                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
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179 TGGCCTGGGAGAACCACCATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGG 238
                            304 GTGACTGTCCGAGCCCAGGGCGGGATACAGTCTGCAGGGAGTGTGAAAAGGGCACCTTTA 363
                                                          119 CCCCTTGCACGGAGCCCTGCGCAACTCCACCTGCCTTGTGTGTCCCCAAGACACCTTCT 178
                                                                                          244 TCCATTCTAAGAACAATTCCATCTGCTGCACCAAGTGCCACAAAGGAACCTACTTGGTGA 303
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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                                                                                                                              59
                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 33
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CLASSIFICATION:
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ZIP: 77210-4433
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Bazzoni, Flavia M.
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Pred. No. 0
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                                                                                                                                                           Mismatches 116;
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; LOCATION: 1.1345
OTHER INFORMATION: /note= "product = Zea mays AP1'
US-08-592-214A-7
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Trocal Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-
TELECOMMUNICATION INFORMATION:
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                   FEATURE:
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                                                                        520 GCTGTCTGTGGCTGGAGGCAGATGTTCTGGG
                                                                                                                                                                                                                                                                                                                                               FEATURE:
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CORRESPONDENCE ADDRESS:
                                                        840 ATTGTCAACGGCGGGAAGCAGATGTTGTGTG 810
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                                                                                                                                 460 GGCGATGGCTGTGCTGCCCCACGAGCACCCTGGGGGAGCTGTCCAGAGCGCTGTGCC 519
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completed: April
me : 142 secs
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TOPOLOGY: lir
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ENTION: Cauliflower Floral Meristem Identify
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OM nucleic - nucleic search, using sw model
                                                                                                   GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
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Run on: April 6, 2003, 13:28:53; Search time 236 Seconds (without alignments) 5296.013 Million cell updates/sec

Title: Perfect score: US-09-993-234-9\_COPY\_146\_700 555

Sequence: 1 CTGGGGGCCCGGGCCCAGGG.........GGCAGATGTTCTGGGTCCAG 555

Scoring table:

IDENTITY\_NUC
Gapop 10.0 , Gapext 1.0

Searched:

Total number of hits satisfying chosen parameters: 4370478

2185239 seqs, 1125999159 residues

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

SIDS2/gcgdata/geneseq/geneseqn-embl/NA1986.DAT: \*
/SIDS2/gcgdata/geneseq/geneseqn-embl/NA1987.DAT: \*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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AAT09420	ABK40265	AAC91477	AAL47186	AAH27782	AAT91180	AAC68777	AAX00925	AAT89427	ID
Death domain conta	cDNA encoding huma	Human PRO779 cDNA.	Human rheumatoid a	Human genomic DNA	Human apoptosis pr	Human death domain	Death domain conta	Death domain conta	Description

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#### ALIGNMENTS

ME SCI INC.	(HUMA-) HUMAN GENOME		
96US-0013285.	12-MAR-1996; 96		
96WO-US16849.			
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	W09/33904-A1.		
/*tag= b			
731251	mat_peptide 7		
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172	sig_peptide 1		
Location/Qualifiers			
	Homo sapiens.		
ds.	inflammation; NF-kappaB;		
containing receptor; DR3; human; apoptosis;	Death domain cont		
ining receptor DR3 cDNA.	Death domain containing receptor		
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This cDNA clone codes for human death domain containing receptor DR3 (see AAW31517), a novel member of the tumour necrosis factor receptor family. It was isolated from a HUVEC cDNA library. Related death domain containing receptor DR3-V1 cDNA (see AAT89426) was isolated from a human testis tumour cDNA library. The genes have also been identified in cDNA libraries of foetal liver, for the containing testing the containing testis to the containing testis tumour cDNA library.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Death domain containing receptor polypeptide(s) DR3 and DR3-V1 - for activation of apoptosis and NF-kappaB, antagonists can be used to treat inflammatory diseases
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           ATGTTCTGGGTCCAG
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ATGTTCTGGGTCCAG
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12-MAR-1996;
17-OCT-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention provides nucleotide sequences encoding death domain containing receptor polypeptides DR3-VI and DR3. The DR3-VI cDNA clone is contained in ATCC deposition No. 97456 and the DR3 cDNA clone is contained in ATCC deposition No. 97757. Recombinant vectors comprising the nucleic acid sequences and optionally the leader sequences are used for the recombinant production of the proteins. The present sequence represents a cDNA encoding the death domain containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 6; Fig 3; 50pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
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(UNMI) UNIV MICHIGAN.
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                       GCCTGGGAGAACCACCATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGCC
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Similarity
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96US-0013285.
96US-0028711.
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1..1254
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Pred. No. 2.7e-144;
; Mismatches 0;
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 The present invention death domain containing
                                       Treating graft-versus-host disease, cancer, immunodeficiency autoimmune disease comprising administering an antibody to De Containing Receptor proteins and a second therapeutic agent
                                                                                                                                                (HUMA-)
(UNMI)
(YUGG/)
                                                                                                                                                                                                                                                                                Human; death domain containing receptor; DR3; cancer; autoimmune disorder; inflammation; cardiovascular disorder; infection;
                                                                                                                                                                                                                                                                                                                                                          AAC68777 standard;
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                        Example 2B; Fig 2;
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28-MAY-1999;
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DB; AAB36265.
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) DILLON P J.
) DIXIT V M.
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) YU G.
) NI J.
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                                                                                                                                                                 HUMAN GENOME SCI INC
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invention provides the containing receptors,
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99US-0136741.
                        273pp; English.
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sequences for DR3-V1. These
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Matches 555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                receptors are involved in apoptosis, and the sequences given can be in the treatment of cancers, infections, cardiovascular disorders arrhythmias, isofhaemia, aneurysms, arterial occlusive diseases, end congenital heart defects, neurodegenerative diseases including alzheimer's and Parkinson's diseases, autoimmune disease such as musclerosis, arthritis, diabetes, Graves' disease, asthma and psorias and to promote angiogenesis and wound healing.
                                                                                                                               AAT91180 standard;
                                           Homo
                                                          Apo-3;
                                                                            Human
                                                                                             14-APR-1998
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                                                                           apoptosis
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555; Conserv
                                                          apoptosis;
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/*tag=
89..160
                                                                           protein Apo-3
                 Location/Qualifiers 89..1342
                                                          human;
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                                                          therapy; drug
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Pred. No. 2.7e-144;
; Mismatches 0;
                                                                                                                               ВP
                                                                           cDNA clone FH20.57
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                                                          screening;
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                                                                                                                                                                                                                                                                                                                                                                CDNA clone FH20.57 (ATCC 55820) codes for a novel polypeptide (see W26709), designated Apo-3, that stimulates or induces apoptotic activity in mammalian cells. It was isolated from a human foetal heart cDNA library by screening with probes (see T91183-84) based on an EST sequence (GenBank locus W71984) that showed homology to the intracellular domain of human TNFR1 and CD95. Amino acid residues 1-181 of Apo-3 are identical to another novel apoptosis polypeptide, Apo-21I (see W26708). Nucleic acids encoding Apo-3 can be used diagnostically for tissue-specific typing and to produce recombinant Apo-3 polypeptides, especially the extracellular domain (amino acids 1-198) or death domain transgenic animals containing cells that express Apo-3 nucleic acid, and knockout animals containing cells that have an altered Apo-3 gene, can be used in drug screening and development.
                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                            Sequence 1634 BP; 300 A; 528 C;
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01-APR-1996;
    446
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hes 555;
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            GGCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCCTTCTACTGCCAA
GGCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGCCAA
                                                TCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCCA
                                                              TCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCCA
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96US-0625328
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/note= "determined by hydropathy analysis"
161..1339
/*tag= c
                                                                                                                                                                                                                                                                                                                       100.0%;
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                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                         Score 555; DB 18;
Pred. No. 2.9e-144;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                             519
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                                     Best Loc
Matches
                                                      Query Match
                                                                                             This invention relates to a method of diagnosing chronic rheumatoid arthritis by digesting human genomic DNA with EcoRI and hybridising with a probe containing a fragment of the present sequence which represents DNA encoding a transmembrane protein. The method is used the diagnosis of chronic rheumatoid arthritis, and for developing ne
                                                                       Sequence 1634 BP;
                                                                                                                                                   Claim 1; Page 14-18; 21pp; Japanese.
                                                                                                                                                                            Diagnosing rheumatoid arthritis by probing digested or comparing expression of mRNA or polypeptide of a
                                                                                                                                                                      transmembrane
                                                                                                                                                                                                                                                                                                                                                        CDS
                                                                                                                                                                                                                                Shiozawa
                                                                                                                                                                                                                                                                  01-NOV-1999;
                                                                                                                                                                                                                                                                                  01-NOV-2000; 2000WO-JP07690
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    146
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DB; AAB97370.
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                                      Conservative
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/product= "Rheumatoid
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                                  Score 555; DB 22;
Pred. No. 2.9e-144;
; Mismatches 0;
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27-MAR-2001;
30-MAR-2001;
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2001JP-0090546.
2001JP-0099990.
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P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to the human DR3 gene, which is associat with rheumatoid arthritis. Certain mutations in the gene can be linked the disease. The sequences can be used to evaluate disease onset and i possibility and to provide therapy and remedies. The present sequence a coding sequence described in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Genomes, particularly DR3 genomic DNA, participating arthritis via mutation, useful in evaluating disease possibility and providing therapy and remedies -
                    Human PRO779
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
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DB; AAO17879.
                                                                                                                                                      ATGTTCTGGGTCCAG
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ilarity 100.0%;
Conservative (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 555; DB 24;
Pred. No. 2.9e-144;
; Mismatches 0;
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be linked to
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Human;

PRO;

antiinflammatory; dermatological;

immunosuppressive; antithyroid;

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The present sequence is one of thirty three nucleic acids encoding PRO polypeptides. The PRO polypeptides, anti-PRO antibodies, agonists and a midgonists are useful for treating and diagnosing immune related disorders such as systemic lupus erythematosus, rheumatoid arthritis, costeoarthritis, juvenile chronic arthritis, spondyloarthropathies, systemic selerosis, idiopathic inflammatory myopathies, Sjogren's systemic vasculitis, sarcoidosis, autoimmune haemolytic anaemia, autoimmune thrombocytopaenia, thyroiditis, diabetes mellitus, commune-mediated renal disease, demyelinating diseases of the central and peripheral nervous systems (such as multiple sclerosis, idiopathic demyelinating polyneuropathy), hepatobiliary diseases of inflammatory demyelinating polyneuropathy), hepatobiliary diseases of conflammatory bowel disease, gluten-sensitive enteropathy and Whipple's disease, autoimmune or immune-mediated skin diseases (such as bullous skin diseases, erythema multiforme, contact dermatitis, psoriasis), allerance sinch as easterna allerance reintitis, ascoriasis;
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15-SEP-1999
15-SEP-1999
29-OCT-1999
30-NOV-1999
01-DEC-1999
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05-JAN-2000
06-JAN-2000
11-FEB-2000
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       allergic
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Hebert C,
Wood WI;
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antidiabetic; n
antiallergic; a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Thirty three nucleic acids encoding PRO polypeptides which are useful in the diagnosis and treatment of immune related disorders, e.g. systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 48; Fig
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DB; AAB50918.
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Henzel W,
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99US-0144732

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99WO-US20111

99WO-US20111

99WO-US21547

99WO-US28313

99WO-US2833

99WO-US30911

2000WO-US30911

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2000WO-US034341

2000WO-US043442

2000WO-US04444

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                                                                                                                                                                                                                                                                                                                                                                                                                                                           33; 218pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             diabetes mellitus
       such
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Kabakoff
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       allergic
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Shelton
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, Tumas D, Watanabe
is, psoriasis),
atopic dermatitis
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Homo

sapiens

WO200153486-A1

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ABK40265
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Best Local
                                                                     gene
                                                                           Human; PRO; benign tumour; malleukaemia; neuronal disorder; immune
                                                                                                                                                                                  ABK40265 standard; cDNA; 1662
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1662 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                food hypersensitivity and urticaria), immunological diseases of the lung (such as eosinophilic pneumonias, idiopathic pulmonary fibrosis and hypersensitivity pneumonitis), transplantation associated disease including graft rejection and graft-versus-host diseases.
                                                                                                                                                                                                                                                             541
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                                                                                                                       encoding
                                                                    therapy;
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                                                                                                                                                                                                                                                   ATGTTCTGGGTCCAG
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llarity 100.0%;
Conservative (
                                                                                                                                            (first
                                                                                                                       human PRO779
                                                                    cytostatic;
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                                                                                                                                           entry)
                                                                          our; malignant tumour; lymphoid malignancy;
sorder; stromal disorder; blastocoelic diso
immune disorder; anglogenic disorder;
                                                                    neuroprotective;
                                                                                                                     polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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Pred. No. 2.9
D; Mismatches
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No. 2.9e-144;
0;
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Best Local S
Matches 555
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30-NOV-1999;
01-DEC-1999;
01-DEC-1999;
05-JAN-2000;
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28-JUL-1999;
17-AUG-1999;
31-AUG-1999;
01-SEP-1999;
                                                                                                                                                                                                                                                                                                                              polypeptides and the polynucleotide sequences encoding them. The PRO polypeptides, agonists, antagonists or anti-PRO antibodies are useful for treating benign or malignant tumours (e.g. renal, kidney, bladder, breast, etc), leukaemias and lymphoid malignancies, other disorders such as neuronal, glial, astrocytal, hypothalamic, glandular, macrophagal, stromal and blastocoelic disorders, inflammatory, immune and angiogenic disorders. The polynucleotide sequences are also useful in gene therapy. ABK40254-ABK40288 encode for the human PRO polypeptides of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Thirty five nucleic acids encoding PRO polypeptides, useful for treating benign or malignant tumours, leukaemias and lymphoid malignancies, inflammatory, angiogenic and immunologic disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                         The
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 50; Fig 23; 302pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GETH
                                                                                                                                                                                                                                                                                                          Sequence 1662 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ashkenazi AJ,
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20-JUL-1999;
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                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                       present invention relates to the isolation of novel human
                                                                                GCCTGGGAGAACCACATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGCC
                                                                                                                              CCTTGCACGGAGCCCTGCGGCAACTCCACCTGCCTTGTGTGTCTCCCCAAGACACCTTCTTG
                                                                                                                                                                                                              CTGGGGGCCCGGGCCAGGCGCACTCGTAGCCCCAGGTGTGACTGTGCCGGTGACTTC
GGCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGCCAA
                                   TCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCCTGTAAGCCA
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                       TCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGGTGTGGCTGTAAGCCA
                                                                    GCCTGGGAGAACCACCATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGCC
                                                                                                                  CCTTGCACGGAGCCCTGCGGCAACTCCACCTGCCTTGTGTGTCCCCAAGACACCTTCTTG
                                                                                                                                                                                                                                                             555;
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99WO-US28313.
99WO-US28301.
99WO-US28634.
2000WO-US00219.
                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pan J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  9908-133459p.
9900-0125252.
9908-140650p.
9908-140653p.
9908-144758p.
9908-146222p.
9908-146222p.
9908-149395p.
9908-149395p.
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99US-123972P.
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i RM, Roy MA,
                                                                                                                                                                                                                                                            0;
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Pred. No. 2.9e-144;
                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                          G; 289 T;
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Smith
                                                                                                                                                                                                                                                                                                          0 other;
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                                                                                                                                                                                                                                                            Indels
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encoding

full-length

or mature

DR3-V1

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RESULT 9
AAT89426
This cDNA clone, deposited as ATCC 97456, codes for human death domain containing receptor DR3-V1 (see AAW31516), a novel member of the tumour necrosis factor receptor family. It was isolated from a cDNA library derived from cells of a human testis tumour. Related death domain containing receptor DR3 cDNA (see AAT89427) was isolated from a HUVEC cDNA library. The genes have also been identified in cDNA libraries of foetal liver, foetal brain, tonsil and leukocyte.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CDS
                                                                                                                                                                 Death domain containing receptor for activation of apoptosis and N
                                                                                                                                                                                                                                                                                   (HUMA-)
                                                                                                                                                                                                                                                                                                                                                           17-OCT-1996;
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                                                                                                                     Claim 2; Page 71-73; 108pp; English.
                                                                                                                                                                                                             WPI; 1997-470812/43.
P-PSDB; AAW31516.
                                                                                                                                                                                                                                                        Dillon PJ,
                                                                                                                                                                                                                                                                                                                                 12-MAR-1996;
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                                                                                                                                                   to treat inflammatory diseases
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                                                                                                                                                                                                                                                                                                    HUMAN GENOME SCI INC
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                                                                                                                                                                                                                                                      Dixit VM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              containing; NF-kappaB;
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304..1481
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198..302
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NF-kappaB, antagonists can be
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sig_peptide
                                                                                               Death
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 produce recombinant polypeptides in transformed host cells. These polypeptides can be used to treat diseases and disorders associated with the inhibition of apoptosis. Antagonists, such as antibodies raised against DR3 VI, can be used to treat diseases and disorders associated with increased apoptosis and for treating inflammatory
                                                                            Death
                                                                                                                25-MAR-1999
                                                                                                                                                     AAX00924
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                                                                                                                                                    standard; cDNA;
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                                                                          receptor; DR3-V1;
/*tag= a
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        "Death domain containing
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                                                                                           polypeptide (DR3-V1)
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12-MAR-1996;
17-OCT-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              "The invention provides nucleotide sequences encoding death domain containing receptor polypeptides DR3-V1 and DR3. The DR3-V1 cDNA clone is contained in ATCC deposition No. 97456 and the DR3 cDNA clone is contained in ATCC deposition No. 97757. Recombinant vectors comprising the nucleic acid sequences and optionally the leader sequences are used for the recombinant production of the proteins. The present sequence represents a cDNA encoding the death domain containing
                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 2;
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P-PSDB;
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(UNMI ) UNIV MICHIGAN.
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ACTGACTGTGGGACCTGCCTGCCTGGCTTCTATGAACATGGCGATGGCTGCGTGTCCTGC
                                                                GGCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGCCAA
                                                                                                                                                        TCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCCA
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                                                                                                                                                                                                                                                              CCTTGCACGGAGCCCTGCGGCAACTCCACCTGCCTTGTGTGTCTCCCCAAGACACCTTCTTG
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                     ACTGACTGTGGGACCTGCCTGGCTTCTATGAACATGGCGATGGCTGCGTGTCCTGC
                                                                                                                                                                                                                                                                                             CACAAGAAGATTGGTCTGTTTTGTTGCAGAGGCTGCCCAGCGGGGCACTACCTGAAGGCC
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                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 555; DB 20; ilarity 100.0%; Pred. No. 3e-144; Conservative 0; Mismatches 0;
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96US-0013285.
96US-0028711.
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/*tag= c
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                                                                                                                                   Query Match
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Matches 555
                                                                                                                                                                                                                                                                              The present invention provides the protein and coding sequences for two death domain containing receptors, designated DR3 and DR3-V1. These receptors are involved in apoptosis, and the sequences given can be used in the treatment of cancers, infections, cardiovascular disorders such as arrhythmias, ischaemia, aneurysms, arterial occlusive diseases, embolisms and congenital heart defects, neurodegenerative diseases including Alzheimer's and Parkinson's diseases, autoimmune disease such as multiple solerosis, arthritis, diabetes, Graves' disease, asthma and psoriasis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Treating graft-versus-host disease, cancer, immunodeficiency or autoimmune disease comprising administering an antibody to Death Containing Receptor proteins and a second therapeutic agent -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; death domain containing receptor; DR3-V1; cancer; autoimmune disorder; inflammation; cardiovascular disorder; infection;
                                                                                                                                                                                                                         Sequence 1783 BP;
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28-MAY-1999;
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)B; AAB36264.
CACAAGAAGATTGGTCTGTTTTGTTGCAGAGGCTGCCCAGCGGGGCACTACCTGAAGGCC
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                                                                                                                                                                                                                                                                  promote angiogenesis and wound healing.
                                                                                                                                                          Similarity
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99US-0136741.
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                                                                                                                                 Score 555; DB 21
Pred. No. 3e-144;
; Mismatches 0
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                                                                                                                                                                                                                      G; 327 T; 0 other;
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                WPI; 1998-240077/21.
P-PSDB; AAW57045.
                                                                         04-OCT-1996;
                                                                                         03-OCT-1997;
DNA encoding apoptosis inducing receptor - which is
                                                                                                                                                                                                          Apoptosis inducing receptor; AIR protein; Type I transmembrane protein; tumour cell
                                                                                                                                                                                                                                 Human apoptosis
                                                                                                                                                                                                                                                                    AAV28700;
                                                                                                                                                                                                                                                                                   AAV28700 standard; cDNA; 1847
                                          Perkins
                                                        (IMMV ) IMMUNEX
                                                                                                                          WO9814565-A1
                                                                                                                                                                                 Homo sapiens
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                                                                                         97WO-US17876
                                                         CORP
                                                                                                                                                        Location/Qualifiers 236..1489
                                                                                                                                                                                                                                 inducing receptor coding sequence
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                                                                                                                                                  /*tag=
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                                                                                                                                                                                                                                                                                   ВP
                                                                                                                                                                                                         human; cell death regulator;
death; autoimmune disease;
 Type
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RESULT 13
ABL64119
ID ABL64
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AC ABL64
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DT 15-MA
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DT 15-MA
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MW Human
KW stoma
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  stomach;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention. AIR is a Type I transmembrane protein, soluble forms of which can be used to regulate cell death in a therapeutic setting. Soluble AIR can also be used in vitro to block apoptosis or AIR-expressing cells, or to screen agonists or antagonists of AIR activity. The cytoplasmic domain of AIR can be used to develop assays for inhibitors of AIR-induced cell death, which is useful to regulate cell death in a therapeutic setting as well as in vitro. Agonists of AIR activity can be used to kill tumour cells that express AIR, or T cells expressing AIR in autoimmune diseases.
           Human;
                          Breast cancer
                                          15-MAY-2002
                                                           ABL64119;
                                                                            ABL64119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 1847
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           cancer;
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                                                                            standard;
  lung;
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                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP;
                         related gene sequence
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  prostate;
         colon; breast; ovary; oesophagus; kidney;
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  pancreas;
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Pred. No. 3e-144;
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  carcinoma;
                         SEQ ID NO: 2456
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  antitumour;
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(AVAL-) AVALON
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05-JUN-2000;
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gene; ds.
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                              2000US-237173P.
2000US-237173P.
2000US-237278P.
2000US-237294P.
2000US-237295P.
2000US-237316P.
2000US-23758P.
2000US-23758P.
2000US-237604P.
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2000US-245084P.
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2000US-236032P
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2000US-236034P
2000US-236101P
2000US-236111P
2000US-236842P
2000US-236891P
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2000US-235711P.
2000US-235720P.
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PHARM.
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PE, DR, 2002-188264/24. Augustus Weaver 2

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Screening for a.... chemical agent to be to 1; for anti-neoplastic agent involves exposing cells to agent to be tested for anti-neoplastic activity, and ng a change in expression of a gene of a signature ger IJ 44pp; gene Q1

set

English

The present invention describes a method (M1) for screening fanti-neoplastic agent. The method involves exposing cells to agent to be tested for anti-neoplastic activity, determining expression of at least one gene (I) of a signature gene set, comprises a sequence (S) selected from 8447 sequences (given comprises a sequence to ABL70110), or is a least (S) for a chemical a change i 'n ABL61664 in

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AAF83770
ID AAF8
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AC AAF8
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         TR3; cell proliferation; leukemia; immunosuppressive; cytostatic; dermatological; antiarthritic; antidiabetic; neuroprotective; car antithyroid; antiinflammatory; antiallergic; T-cell-inhibitor; ds
                                                                                             06-AUG-2001
                                                                  Nucleotide
                                                                                                                                                 AAF83770 standard;
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                                                                  of human TR3
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99.1%;
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2.3e-140;
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            neuroprotective; cardiant;
T-cell-inhibitor; ds.
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                                                                                                                                                                                                   The invention relates to a composition comprising a biologically active PR3-specific binding agent (I) that binds to TR3 and inhibits the proliferation of cells expressing TR3. (I) identified by the methods are useful for treating a subject suspected of having a disease associated with a proliferation of cells expressing TR3 especially leukemias or lymphomas or a T-cell mediated disease especially autoimmune diseases such as myasthenia gravis, systemic lupus erythamatosus, rheumatoid arthritis, diabetes, multiple sclerosis, sarcoidosis, myocarditis, thyroiditis and tumours. (I) is also useful for treating a subject suspected of having graft-versus-host disease, rejection of a transplanted organ such as heart, liver, lung, kidney, pancreas, bowel, skin or an appendage, or inflammatory diseases, allergies and contact dermatitis. The present sequence represents the nucleotide sequence of
                                                                                                                                                                                                                                                                                                                                                                                                                                     Composition for treatment of T-cell mediated disease cancer comprises a biologically active TR3-specific especially a monoclonal antibody -
                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 72;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                P-PSDB; AAB84941.
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(WEGM/) WEGMANN K W.
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CACAAGAAGATTGGTCTGTTTTGTTGCAGAGGCTGCCCAGCGGGGCACTACCTGAAGGCC
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deletion which alters the reading frame"
/transl_except= "(pos:558..559, as:Cys)"
/note= "this codon has an apparent 1 nucleotide
deletion which alters the reading frame"
/transl_except= "(pos:638..639, as:Leu)"
/note= "this codon has an apparent 1 nucleotide
deletion which alters the reading frame"
/transl_except= "(pos:718..719, as:Met)"
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27-MAR-2001;
30-MAR-2001;
      arthritis
                        Genomes,
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SHIOZAWA S.
  particularly DR3 genomic via mutation, useful in
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Best Local Similarity
Matches 487; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to the human DR3 gene, which is associat with rheumatoid arthritis. Certain mutations in the gene can be linked the disease. The sequences can be used to evaluate disease onset and possibility and to provide therapy and remedies. The present sequence a coding sequence described in the exemplification of the invention.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

### SUMMARIES

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## ALIGNMENTS

AUTHORS TITLE	REFERENCE		ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION		DEFINITION	LOCUS	BM695193	RESULT 1
Bonaldo,M.F., Lennon,G. and Soares,M.B. Normalization and subtraction: two approaches to facilitate gene	Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 510)	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;	Homo sapiens	human.	EST.	BM695193.1 GI:19008451	BM695193	UI-E-CQ1-aev-i-03-0-UI 5', mRNA sequence.	UI-E-CQ1-aev-i-03-0-UI.rl UI-E-CQ1 Homo sapiens cDNA clone	BM695193 510 bp mRNA linear EST 28-FEB-2002		

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                                                                                                                                                                      LeuLysArgTrpArgGlnGlnGlnProAlaGlyLeuGlyAlaValTyrAlaAlaLeuGlu 60
                                                                       CGCATGGGCTGGACGGCTGCGTGGAAGACTTG 216
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Genome Res.
97044477
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Tissue Procurement: Dr. Gregg Hageman
cDNA Library preparation: Dr. M. Bento Soares, Univeristy of Iowa
cDNA Library Arrayed by: Dr. M. Bento Soares, Univeristy of Iowa
DNA Sequencing by: Dr. M. Bento Soares, Univeristy of Iowa
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451 Eckstein Medic
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Program for Rat Gene Discovery and Mapping
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /lab_host="DH10B (Life Technologies) (T1 phage resistant)"
/note="Organ: eye; Vector: pT7T3-Pac (Pharmacia) with a
modified polylinker; Site_1: ECOR I; Site_2: Not I;
UT-E-CQ1 is a normalized CDNA library containing the
following tissue(s): optic nerve. The library was
constructed according to Bonaldo, Lennon and Soares,
Genome Research, 6:791-806, 1996. First strand cDNA
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/dev_stage="adult"
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/clone_lib="UI-E-CQ1"
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US-09-993-234-6_COPY_339_409 (1-71) x BM666370 (1-550)
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Program for Rat Gene Discovery and Mapping
University of Iowa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Genetics (www.resgen.com).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   451 Eckstein Medical Research Building Iowa City, IA 52242, USA
Tel: 319 335 8250
Fax: 319 335 9565
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Bonaldo, M.F., Lenno
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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CDNA Library preparation: Dr. M. Bento Soares, Univeristy of Iowa
CDNA Library Arrayed by: Dr. M. Bento Soares, Univeristy of Iowa
DNA Sequencing by: Dr. M. Bento Soares, Univeristy of Iowa
Clone Distribution: Researchers may obtain clones from Research
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/note="Organ: eye; Vector: pT7T3-Pac (pharmacia) with a
modified polylinker; Site_1: EcoR I; Site_2: Not I;
UI-E-QQ1 is a normalized cDNA library containing the
following tissue(s): optic nerve. The library was
constructed according to Bonaldo, Lennon and Soares,
Genome Research, 6:791-806, 1996. First strand cDNA
synthesis was primed with an oligo-dT primer containing a
Not I site. Double stranded cDNA was ligated to an EcoR I
adaptor, digested with Not I, and cloned directionally
into pT7T3-Pac vector. The oligonucleotide used to prime
the synthesis of first-strand cDNA contains a library tag
sequence that is located between the Not I site and the
                                                                                                                                                                                                                                                                             TAG_TISSUE=human optic nerve
TAG_SEQ=CCATTAAGTG*
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/clone_lib="UI-E-CQ1"
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similar to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nlh.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,
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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
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1 (bases 1 to 687)
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BE670189
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              info@image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cDNA Library Preparation: M. Bento Soares, Ph.D.
CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL, send email to:
                                                                                                                                                              111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        quality sequence stop: 466.
                                                                                                                                                                                                                                /note-*Organ: lung; Vector: pT7T3D-Pac (Pharmacia) with modified polylinker; Plasmid DNA from the normalized library NCI_CGAP_LUS was prepared, and ss circles were made in vitro. Pollowing HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from a pool of 5,000 library was PCR-amplified cDNAs from a pool of 5,000 library was PCR-amplified cDNAs from a pool of 5,000 library was PCR-amplified cDNAs from a pool of 5,000 library was PCR-amplified cDNAs from a pool of 5,000 library was PCR-amplified cDNAs from a pool of 5,000 library was PCR-amplified cDNAs from a pool of 5,000 library was PCR-amplified cDNAs from a pool of 5,000 library was PCR-amplified cDNAs from a pool of 5,000 library was PCR-amplified cDNAs from a pool of 5,000 library was properties.
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                                                                                                                                                                                              clones made from the same library (cloneIDs 1414920-1417991 and 1520904-1522439). Subtraction by Bento
                                                                                                                                                                                                                                                                                                                                                                                      /clone="IMAGE:3284086"
/clone_lib="NCI_CGAP_Lu24"
                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                    /tissue_type="carcinoid"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                      /lab_host="DH10B"
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SW:WSL1_HUMAN
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  Length:
Matches:
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                  Best Local Similarity:
Query Match:
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AUTHORS
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VERSION
KEYWORDS
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AW074008/c
      DB:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CTCAAGCGCTGGCGCAGCAGCAGCCGGGGCCTCGGAGCCGTTTACGCGGCCCTGGAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution
Clone distribution: NCI-CGAP clone distribution
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EST
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AW074008.1
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Technologies, Inc. cDNA Library Arrayed by: Christa Prange, Th
I.M.A.G.E. Consortium DNA Sequencing by: Washington University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tissue Procurement: Chris Moskaluk, M.D., Ph.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: cgapbs-r@mail.nih.gov
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National Cancer Institute, Cancer Genome Anatomy
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1 (bases 1 to 618)
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Location/Qualifiers
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Site_1: Sall; Site_2: NotI;
Primer: Oligo dT. Library
Technologies."
                                                                                                                                                                                                                                                                                                        /clone="IMAGE:2575504"
/clone_lib="NCI_CGAP_GU1"
/tlssue_type="2 pooled high-grade transitional cell
                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                      /lab_host="DH10B"
1.8e-37
361.00
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                                                                                                                                                                                                                                                                                                                                                                                                               Email: cgapbs=r@mail.nih.gov
Tissue Procurement: Ash Alizadeh, John Byrd, M.D., Mike Grever,
M.D., Louis M. Staudt, M.D., Ph.D.
cDNA Library Preparation: M. Bento Soares, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
cDNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 1237 Std Error: 0.00
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tg18c01.x1 NCI_CGAP_CLL1 Homo sapiens cDNA clone IMAGE:2109120
similar to TR:000278 000278 LYMPHOCYTE ASSOCIATED RECEPTOR OF I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D.
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 587)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (1997)
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Library is normalized, and was constructed
Soares and M.Fatina: Bonaldo."
192 c 176 g 120 t
                                                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:2109120"
                                                                                                                                                                                                                                                                   /clone_lib="NCI_CGAP_CLL1"
/tissue_type="B-cell, chronic lymphotic leukemia"
                                                                                                                                                                                                                                                        /lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
        1.4e-36
354.00
                                                                                                             double-stranded cDNA was ligated to Eco RI (Pharmacia), digested with Not I and cloned into I and Eco RI sites of the modified pT7T3 vector.

of a normalized, and was constructed by Bento
        Length:
Matches:
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        587
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AI811528/c
LOCUS
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AUTHORS
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Seq primer: -40UP from Gibco
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CDNA Library Preparation: Life Technologies, Inc
CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome
Clone distribution: NCI-GGAP clone distribution
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               High quality sequence stop: 392.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tissue Procurement: Christopher Moskaluk, M.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
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1 (bases 1 to 623)
                                                                                                                                                       Average insert size 1.75 kb. 11538-014"
                                                                                                                                                                                                    /note="Organ: uterus; Vector: pCMV-SPORT6; Site_1: Sall;
Site_2: Not1; Cloned unidirectionally. Primer: Oligo dT
                                                                                                                                                                                                                                                 /clone="IMAGE:2262491"
/clone_lib="NCI_CGAP_Ut1"
/clone_tib="NCI_CGAP_Ut1"
/tissue_type="well-differentiated endometrial
adenocarcinoma, 7 pooled tumors"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
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Sequence evaluation of four pooled-tissue normalized bovine libraries and construction of a gene index for cattle Genome Res. 11 (4), 626-630 (2001)
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279811 MARC 3BOV Bos
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EST.
                                                                                                                                                                                                                                                                                                                        Email: smith@email.marc.usda.gov
Single pass sequencing. Bases called and alt_trimmed with phred
v0.980904 e. Vector identified by cross_match with the -minscor
                                                                                                                                                                                                                                                                                                                                                                                                               USDA, ARS, US Meat Animal Research Center
PO Box 166, Clay Center, NE 68933-0166, USA
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Fax: 402 762 4390
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Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora;
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          /note="Vector: pCMV SPORT6; Site_1: Not1; Site_2: Sal1; Library made from pooled tissue from marrow, alveolar macrophage, ovary, fetal semitendonosus muscle, and fetilongissimus muscle."

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                                                                                                                                 /organism="Bos taurus"
/db_xref="taxon:9913"
/clone_lib="MARC 3BOV"
                                                                                                 /tissue_type="pooled"
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                                                                                                                                                                                                                                                                                                                                                                                                                            This clone is available royalty-free through LINL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Seq primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      xp99a04.x1 Soares_NHCeC_cervical_tumor Homo sapiens cDNA clone IMAGE:2748462 3' similar to TR:000278 000278 LYMPHOCYTE ASSOCIATED RECEPTOR OF DEATH 7. [2] TR:000279 ; mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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National Cancer Institute, Cancer Genome Anatomy
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                                                                                       Q)
                                                                                    T 3']; double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is normalized; constructed by Bento Soares and M.Fatima Bonaldo."

a 189 c 178 g 115 t 1 others
                                                                                                                                                                                   /clone_lib="Soares_NHCeC_cervical_tumor"
                                                                                                                                                                                                                                                                                                                       /clone="IMAGE:2748462"
                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
/db_xref="taxon:9606"
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Conservative:
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                   Email: yongsung@mail.kribb.re.kr
Plate: 48 row: G column: 11
High quality sequence stop: 647.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tel: +82-42-860-4470
Fax: +82-42-860-4409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Korea Research Institute of Bioscience & Biotechnology 52 Eoeun-dong Yuseong-gu, Daejeon 305-333, South Korea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 (bases 1 to 647)
Kim, N.S., Hahn, Y.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Kim YS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21C Frontier Korean EST Project
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                 110
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           inote-*Organ: Stomach; Vector: pT7T3-Pac; Site_1: EcoRI; Site_2: NotI; The S22SNU16 library was contributed by the Soares laboratory and it was constructed as described by Bonaldo, M.F., Lennon, G. and Soares, M.B. (1996), Genome Research 6(9): 791-806. RNA was prepared from harvested cells of SNU-16 culture. SNU-16 cell was obtained from Korean Cell Line Bank (KCLB). SNU-16 was established from ascitic fluids of Korean patients by Park J.G. et al. (1990), Cancer Res 50: 2773-2780."
                                                                                                                                                                                                                      /tissue_type="Ascites"
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                                                                                                                                                                                                                                                                                                    /organism="Homo sapiens"
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/clone="S22SNU16n1-48-G11"
/clone_lib="S22SNU16n1"
                                                                                                                                                                                                          /lab_host="DH10B"
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E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,
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                                                                                                                                                                                                                                                                                                                                                         M.D., Louis M. Staudt, M.D., Ph.D.

CDNA Library Preparation: M. Bento Soares, Ph.D.

CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome

Clone distribution: NCI-CGAP clone distribution

found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     544 DP mRNA linear EST 30-MAR-1999 tg17f01.x1 NCI_CGAP_CLL1 Homo sapiens cDNA clone IMAGE:2109049 3' slimilar to TR:000278 000278 LYMPHOCYTE ASSOCIATED RECEPTOR OF DEATH 7. [2] TR:000279 ; mRNA sequence.
                                                                                                                                                                                                                                                                                                      www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 1246 Std Error: 0.00
Seg primer: ~40Up from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Ash Alizadeh, John Byrd, M.D.,
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National Cancer Institute, Cancer Genome Anat
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                                                                                                                                                                                                                                                                       quality sequence stop: 348.
Location/Qualifiers
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T 3']; double-stranded CDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Thrarv is normalized, and was constructed by Bento
                                                                       /organism="Homo sapiens"
/db_xref="taxon:9606"
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/clone="InMagE:2109049"
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/tlssue_type="B-cell, chronic lymphotic leukemia"
/lab_host="DH10B"
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/note-"Organ: Stomach; Vector: pT7T3-Pac; Site_1: EcoRI; S1te_2: NotL; The S2ZSNU16 library was contributed by the Soares laboratory and it was constructed as described by Bonaldo, M.F., Lennon, G. and Soares, M.B. (1996), Genome Research 6(9): 791-806. RNA was prepared from harvested
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/clone="S22SNU16n1-85-D08"
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25 1 (bases 1 to 633)
26 Arakawa,T., Carninci,P., Fukuda,S., Furuno,M., Hanagaki,T., Hara,A., Hiramoto,K., Hori,F., Ishii,Y., Ito,M., Kawai,J., Konno,H., Kouda,M., Koya,S., Matsuyama,T., Miyazaki,A., Nomura,K., Ohno,M., Okazaki,Y., Okido,T., Salto,R., Sakai,C., Sakai,K., Sano,H., Sasaki,D., Shibata,K., Shinagawa,A., Shiraki,T., Sogabe,Y., Suzuki,H., Tagami,M., Tagawa,A., Takahashi,F., Takeda,Y., Tanaka,T., Toya,T., Muramatsu,M. and Hayashizaki,Y.

NIKEN Mouse ESTs (Arakawa,T., et al. 2001)
20 Jun 30, 2000 this sequence version replaced gi:8877385.
Contact: Yoshihide Hayashizaki
Contact: Yoshihide Hayashizaki
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Email: genome-res@gsc.riken.go.jp,

URL:http://genome.gsc.riken.go.jp/

Carninci,P., Shibata,Y., Hayatsu,N., Sugahara,Y., Shibata,K., Itoh
,M., Konno,H., Okazaki,Y., Muramatsu,M. and Hayashizaki,Y.

Normalization and subtraction of cap-trapper-selected cDNAs to
prepare full-length cDNA libraries for rapid discovery of new
genes. Genome Res. . 10 (10), 1617-1630 (2000)

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Watahiki,M., Yoneda,Y., Ishikawa,T., Ozawa,K., Tanaka,T., Matsuura
,S., Kawai,J., Okazaki,Y., Muramatsu,M., Inoue,Y., Kira,A. and
                                                                                                                                                                                                                              Laboratory for Genome Exploration Research Group, RIKEN Genomic Sciences Center(GSC), Yokohama Institute of Physical and Chemical Research (RIKEN) 1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Jappels B1-45-503-9222
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                                                                LeuLysArgTrpArgGlnGlnGlnProAlaGlyLeuGlyAlaValTyrAlaAlaLeuGlu 60
                                                                                                                                   AlaGluIleGluAlaValGluValGluIleGlyArgPheArgAspGlnGlnTyrGluMet 40
                                                                                                                                                                                                                MetAspAlaValProAlaArgArgTrpLysGluPheValArgThrLeuGlyLeuArgGlu 20
                                                                                                                                                                                              ATGGATGCGGTCCCAGCACGAAGGTGGAAGGAGTTCGTGCGCACGCTGGGGCTGCGGGAA 194
                          CTCAAGCGCTGGCGTCAGCAGCAGCCTGCAGGCCTCGGTGCCATCTATCG-GCTCTGGAG
                                                                                                               GCGGAAATTGAAGCCGTGGAGGTGGAAATCTGCCGCTTCCGAGACCAGCAGTATGAGATG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Encyclopedia Project of Genome Exploration Genomic Sciences Center and Genome Science Division of Experimental Animal Research in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   encyclopedia: real-time sequence clustering for construction of a nonredundant cDNA library. Genome Res. 11 (2), 281-289 (2001) Yamanaka, I. Kiyosawa, H., Kondo, S., Saito, T., Shinagawa, A., Aizawa K., Fukuda, S., Hara, A., Itoh, M., Kawai, J., Shibata, K., Arakawa, T., Shil, Y. and Hayashizaki, Y. Mapping of 1903Z mouse cDNAs on mouse chromosomes. J. Struct. Func. Genomics 2 pre, L72-L86 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       prepare mouse tissues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cDNA library was prepared and sequenced in Mouse Genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Please visit our web site (http://genome.gsc.riken.go.jp/) for further details.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequencing pipeline with 10 (11), 1757-1771 (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Computer-based methods for the mouse full-length cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Konno, H., Fukunishi, Y., Shibata, K., Itoh, M., Carninci, P., Sugahara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RIKEN integrated sequence analysis (RISA) system--384-format requencing pipeline with 384 multicapillary sequencer. Genome Res.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    prepared and sequenced in Mouse Genome Encyclopedia
Project of Genome Exploration Research Group in Riken
Genomic Sciences Center and Genome Science Laboratory in
RIKEN. Division of Experimental Animal Research in Riken
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /dev_stage="0 day neonate"
/lab_host="DH10B"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone_lib="RIKEN full-length enriched, 0 day neonate
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Mismatches:
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MEDLINE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: msoares@blue.weeg.uiowa.edu
cDNA Library Preparation: M.B. Soares Lab Clone distribution:
clones will be available through Research Genetics (www.resgen.com)
This clone is also available through the I.M.A.G.E. Consortium at
LLNL (info@image.llnl.gov). IMAGE ID- 1787481
Seq primer: M13 Forward.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Program for Rat Gene Discovery and Mapping University of Iowa 451 Eckstein Medical Research Building Iowa City, IA 52242, USA Tel: 319 335 8250
Fax: 319 335 9565
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UI-R-C2p-oh-b-10-0-UI.rl UI-R-C2p Rattus norvegicus cDNA clone
UI-R-C2p-oh-b-10-0-UI 5', mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Genome Res. 6 (9), 791-806 (1996)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EST.
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                         49
                                                                                                                                      /lab_host="DHIUB (Life Technologies)"
/lab_host="DHIUB (Life Technologies)"
/note="Vector: p773D-Pac (Pharmacia) with a modified
polylinker; Site_1: Not I; Site_2: Eco RI; The UI-R-C2p
library is a subtracted library derived from the UI-R-C1
library, which is a subtracted library derived from the
UI-R-C0 library. The UI-R-C0 library consisted of a
mixture of individually tagged normalized libraries
constructed from rat placenta, adult lung, brain, liver,
kidney, heart, spleen, ovary, muscle, 8, 12 and 18-day
embryo. The tag is a string of 3-5 nucleotides present
between the Not I site and the oligo-dT track which allows
identification of the library of origin of a clone within
the mixture. The subtracted library (UI-R-C2p) was
constructed as follows: PCR amplified CDNA inserts from
UI-R-C1 clones from which 3' ESTs had been derived was
used as a driver in a hybridization with the UI-R-C1
library in the form of single-stranded circles. The
remaining single-stranded circles (subtracted library)
converted to double-stranded circles subtracted dinto DHIUB bacteria (Life Technologies) to generate the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  norvegicus
                                                into DH10B bacteria (Life Technologies) to generate the UI-R-C2p library. This procedure has been previously described (Bonaldo, Lennon and Soares, Genome Research (791-806, 1996)"
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/dev_stage="adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone="UI-R-C2p-oh-b-10-0-UI"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BQ027499 499 bp mRNA UI-H-COO-aqq-a-09-0-UI.sl NCI_CGAP_Sub9 IMAGE:3104895 3', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                CDNA Library preparation: Dr. M. Bento Soares, University of Iowa CDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa DNA Sequencing by: Dr. M. Bento Soares, University of Iowa Clone Distribution: Clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://lmage.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 499)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Dr. Jose Mercuende
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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/note="vector: p7773-Pac (Pharmacia) with a modified polylinker; Site_1: EcoR I; Site_2: Not I; tissues: Cholonic mucosa with Crohns disease, Cholonic mucosa with ulcerative colitis, Fetal thymus, Cervix, Cervical adenosquamous carcinoma, Ligament cells, Prostate carcinoma, Bladder carcinoma, Brain oligodenroga: NCI_CGAP_Sub9 is a subtracted cDNA library constructed according to Bonaldo, Lennon and Soares, Genome Research, 6:791-806, 1996. First strand cDNA synthesis was primed with an oligo-dr primer containing a Not I site. Double stranded cDNA was ligated to an EcoR I adaptor, digested with Not I, and cloned directionally into p7773-Pac vector. The oligonucleotide used to prime the synthesis of first-strand cDNA contains a library tag sequence that is
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/clone_lib="NCI_CGAP_Sub9"
/tissue_type="mixed"
/dev_stage="mixed"
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/db_xref="taxon:9606"
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Tel: 301 838 3528
Fax: 301 838 0208
                                                                                                                                                                                                                                           Contact: John Quackenbush
The Institute for Genomic Research
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                                                                                                                                                       Email: johnq@tigr.org
Plate: 207
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                             Location/Qualifiers
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# SUMMARIES

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2 (bases 1 to 1254)
Chinnaiyan,A.M., O'Rourke,K., Yu,G.-L., Lyons,R.H.,
Duan,R., Xing,L., Gentz,R., Ni,J. and Dixit,V.M.
Direct Submission
                                                                                             Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates; 1 (bases 1 to 1254)
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                                                                                                                               Homo sapiens.
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/organism="unknown"
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Pred. No. 9.6e-133;
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and Dixit, V.M.
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555; Conserv
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/codon_start=1
/codon_start=1
/product="death receptor 3"
/db_xref="g1:1669512"
/db_xref="g1:1669512"
/db_xref="g1:1669512"
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/f1669AGHYLKAPCTEPCGNSTCLVCPODTFLAMENHHNSECARCQACDEQAQV
ALENCSAVADTRCGCKPCMFVECQVSQCYSSSPPYCQPCLDCGALHRHTRLLCSRRDT
DCGTCLPGFTEHEDGGCVSCPTSTIGSCPERCANVCGWRQMFWYQVLLAGLVYPLLIGA
TLTYTYRHCWPHKPLVTADEAGMEALTPPPATHLSPLDSAHTLLAPPDSSEKICTVQL
VGNSWTPGYFETGEALCPQYTWSWDQLPSRALGPAAPTLSPESPAGSPAMMLQPGPQ
LYDVMDAVPARRMKEFVETLGLREAETIEAVEVETGREFRDQQYEMLKRWRQQQPAGLGA
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/db_xref="taxon:9606"
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                                               CCATGCCTAGACTGCGGGGCCCTGCACCGCCACACACGCTACTCTGTTCCCCGCAGAGAT 420
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Degli-Esposti,M.A. and Goodwin,R.G.
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/protein_id="Apa0018.1"
/db_xref="Gd:1778764"
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/translation="MEORPRGCAEVECCAVCOCOTELAWENHHUSECARCQACDEORAOV
ALENCSAVADTRCGCREGMETVECOVSOCVSSSPFYCOPCLDGALHRHTRLLCSRAOV
ALENCSAVADTRCGCREGMETVECOVSOCVSSSPFYCOPCLDGALHRHTRLLCSRAOV
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1 (bases 1 to 1254)
Screaton,G.R., Xu.X.N., Olsen,A.L., Cowper,A.E., Tan,R., McMichael,A.J. and Bell,J.I.

LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing
Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
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Submitted (19-MAR-1997) Molecular Immunology Group, Institute of
Molecular Medicine, John Radcliffe Hospital, Headington, Oxford
                                                               Similarity 100 55; Conservative
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                                                                                                                                                                      /db_xref="G1:2071949"
//translation="MEDRREGGAVAAALLLYLLGARAQGGTRSPRCDCAGDEHKKIG
//translation="MEDRREGGAVAAALLLYLLGARAQGGTRSPRCDCAGDEQASQV
LECCRGCPAGHYLKAPGTEPGGNSTCLVCPQDTFLAWENHHNSECARCQACDEQASQV
ALENCSAVADTRCGCKRCWFVECQVSQCVSSSPFYCOPCLDCGALHRHTRLLCSRRDT
DCGTCLPGFYEHGDGCVSCPTSTLGSCCPERCAAVCGWRQWFWQLLAGLVPPLLGA
TLTYTRHCWPARDAGAGMALTPPPATHLSPLDSAHTLLAPPDSSEKICTVQL
VGNSWTPGYPETQEALCPQVTWSWDQLFSRALGPAAAPTLSPESPAGSFAMMLQPGPQ
LYDVMDAVPARRWKEFVRTLGLREAEIEAVEVEIGRFRDQQYEMLKRWRQQQPAGLGA
                                                                                                                                                VYAALERMGLDGCVEDLRSRLQRGP"
420 c 407 g 226
                                                                                                                                                                                                                                                                                                                                                           /function="mediates apoptosis"
/note="LARD-la; membrane protein;
TNF-R1; contains a death domain"
                                                                                                                                                                                                                                                                                                              /product="lymphocyte associated receptor of death
/protein_id="AAC51306.1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
/db_xref="taxon:9606"
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                                                             Score 555; DB 9;
Pred. No. 9.6e-133;
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LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
                                                                                                                                                                                                                                                                                                                                                                                                  Human lymphocyte associated receptor of death alternatively spliced, complete associated receptor of death reasons are sensitively spliced.
                                                                                                                                              Direct Submission
Submitted (19-MAR-1)
Molecular Medicine,
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                                                                                                                                                                                       Screaton, G.R.
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1 (bases 1 to 1257)
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                                                                   /organism="Homo sapiens"
/db_xref="taxon:9606"
/cell_line="HeLa"
                /function="mediates apoptosis"
/note="LARD-1b; membrane prote
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                                         #11_type="lymphocyte"
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ilarity 100.0%;
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/protein_id="AAC51307.1"
/protein_id="AAC51307.1"
/db_xref="Git:20712951"
/translation="MEQRPRGCAAVAAALLIVILGARAQGGTRSPRCDCAGDEHKKIG
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ALENCSAVADTRCGCKPGMFVECQVSQCVSSSPFYCQPCLDCGALHRHTLLCSRRDT
DCGTCLFGFYEHGDGCVSCPTSTLGSCCERCAAVCGWRQMFWVQVLLAGLVVPLLLGA
TLTYTYEHCHWPHKELVTAADEAGMEALTPPPATHLSPLDSAHTLAPDSSEKICTVQ
LVGNSWTPGYPETQEALCPQVTWSWDQLPSRALGPAAAPTLSPESPACSPAMMIQPGP
QLYDVADAVPAARWKEFVRTLGLREAEIEAVEVEIGRFRDQQYEMLKRWRQQQPAGLG
AVYAALERMGLDGCVEDLRSRLQRGP"
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3' splice site"
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                       mRNA,
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Pred. No. 9.6e-133;
; Mismatches 0;
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ACTGACTGTGGGACCTGCCTGCCTGGCTTCTATGAACATGGCGATGGCTGCGTGTCCTGC
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2 (bases 1 to 1634)
Marsters, S.A., Sheridan, J.P., Donahue, C.J.,
Goddard, A.D., Bauer, K.D. and Ashkenazi, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota;
Mammalia; E
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Marsters, S.A., Sheridan, J.P., Donahue, C.J., Pitti, R.M., Gray, C.L., Goddard, A.D., Bauer, K.D. and Ashkenazi, A.
Apo-3, a new member of the tumor necrosis factor receptor family, contains a death domain and activates apoptosis and NF-kB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted (15-OCT-1996) Molecular Bruno Blvd., South San Francisco,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Direct Submission
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VGNSWTBGYPETQEALCPQVTWSWDQLDSRALGPAAAPTLSPESPAGSPAMMLQBGPQ
LYDWDAVPARRWKEFVTLGLREBETAVEVEIGRFRDQQYEMLKRWRQQQPAGLGA
VYAALERWGLDGCVEDLRSRLQRGP"
a 528 c 519 g 287 t
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LECCRGCPAGHYLKAPCTEPGCNSTCLVGPPDTFLAWENHHUSEGRECAGDCDEQASOV
ALENCSAVADTRGGKRGMFYBCOVSOCVSSSPFYCOPCLDCGALHRTFRLLCSRDTD
DCGTCLPGFYEHGDGCVSCPTSTLGSCPERCAAVCGWROMFWVQVLLAGLVVPPLLLGA
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NF-kB; TNF receptor family member"
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89. .1342
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/protein_id="AAB39714.1"
/db_xref="GI:1763293"
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/db_xref="taxon:9606"
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Genentech, Inc. (US)
Location/Qualifiers
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Ashkenazi, A.J., Bake
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Stone, D.M., Watanabe, C.K. and Wood, W.I.
Compositions and methods for the treatment of tumour
Patent: WO 0153486-A 23 26-JUL-2001;
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Ashkenazi, A.J., Goddard, A.,
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/db_xref="taxon:9606"
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Pitti,R.M., Roy,M.A.,
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                      CCTTGCACGGAGCCCTGCGGCAACTCCACCTGCCTTGTGTGTCCCCAAGACACCTTCTTG
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Pred. No. 9.3e-133;
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TITLE
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Best Local Similarity
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                                                                                                                                           GCCTGGGAGAACCACATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGCC
                                                                                                                                                                                            CACAAGAAGATTGGTCTGTTTTGTTGCAGAGGCTGCCCAGCGGGGCACTACCTGAAGGCC 120
                                                                               TCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCCGCTGTGGCTGTAAGCCA 300
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GGCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGCCAA
                                                              TCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCCA
                                                                                                                                                                                                                                                                                                                                                                                          554;
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Submitted (03-JAN-1997) Molecular Biotechnology, University of
Submitted (1705 NE Pacific Street, HSB-K380, Seattle, WA 98195,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo
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Chaudhary, P.M. and Hood, L.E.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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<1. .1238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /product="death domain receptor 3"
/protein_id="AAB41432.1"
/protein_id="AAB41432.1"
/protein_id="AAB41432.1"
/db_xref="GI:1800293"
/db_xref="GI:1800293"
/translation="GCAVAAALLLVLLGARAQGGTRSPRCDCAGDEQASQVALENCS
CPAGHYLKAPCTEPCGNSTCLVCPQDTFLAWENHHNSECARCQACDEQASQVALENCS
AVADTROGCKPGWFVECQVSGCVSSSPFYCQPCLDCGALHRTRLLCSRRDTDCGTCL
LGFYEHGDGCVSCPTSTLGSCPERCAAVCGWRQMFWVQVLTQLVGATLTYT
RHCWPHKPLVTADEAGMEALTPPPATHLSPLDSAHTLLAPPDSSEKICTVQLVGNSWT
PGYPETQEACKPGVTWSWDQLPSRALGPAAAPTLSPESPAGSPAMMLQPGPQLYDVMD
AVPARBKEFYRTLGLÆAEIEAVEVEIGRFRDQQYEMLKRWRQQQPAGLGAVYAALE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /gene="DDR3"
/function="mediates apoptosis when cross-linked"
/function="mediates apoptosis when cross-linked"
/note="A TWER1-related death-domain containing receptor;
/note="A TWER1-related death-domain containing receptor;
DR3, WSL-LR, Apo-3, TRAMP; transmembrane form; similar to
human EST clone 298913, GenBank Accession Number N71143"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RMGLDGCVEDLRSRLQRGP*
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/db_xref="taxon:9606"
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97205335
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TRAMP, a novel apoptosis mediating receptor with sequence homology to tumor necrosis factor receptor 1 and Fas(Apo-1/CD95)
Immunity 6 (1), 79-88 (1997)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Direct Submission
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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               286
                                                                                  /product="apoptosis-mediating receptor TRAMP"
/protein_id="AACS1192.1"
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TCLLGFYEHGDGCVSCCFTSTLGSCPERCAAVCGWRQMFWQVLLAGLVVPLLLGATLT
YTYRHCWPHKPLVTADEAGMEALTPPPATHLSPLDSAHTLLAPPDSSEKICTYQLVGN
               ALERMGLDGCVEDLRSRLQRGP
                                                  SWTPGYPETQEALCPQVTWSWDQLPSRALGPAAAPTLSPESPAGSPAMMLQPGPQLYD
VMDAVPARRWKEFVRTLGLREAEIEAVEVEIGRFRDQQYEMLKRWRQQQPAGLGAVYA
                                                                                                                                                                                                                                                                                                              /tissue_type="fetal lung"
<1. .1245
                                                                                                                                                                                                                                                                                                                                                                    /organism="Homo sapiens"
/db_xref="taxon:9606"
/chromosome="1"
                                                                                                                                                                                                                                     /codon_start=1
                                                                                                                                                                                                                                                      /function="mediates apoptosis"
/note="contains a death domain;
                                                                                                                                                                                                                                                                                                                                                  /map="1p36.2"
                                                                                                                                                                                                                                                                                                                                                                                                                                         ocation/Qualifiers
                                                                                                                                                                                                                                                                                          function="activates NF-kB"
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Query Match Best Local Similarity

99.7%; 99.8%;

Score Pred.

553.4; DB 9; No. 2.4e-132;

Length 1557

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JOURNAL
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AUTHORS
TITLE
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AUTHORS
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ORGANISM
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                                                                                                                                                                                                                                                Human lymphocyte as spliced, complete of U94510 U94510.1 GI:207196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             554;
                                      Direct Submission
Submitted (19-MAR-1997) Molecular Immunology Group, Institute
Molecular Medicine, John Radcliffe Hospital, Headington, Oxfor
9DU, UK
                                                                                                                        LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
                                                                                                                                                           Screaton, G.R., Xu, X.N., Olsen, A.L., McMichael, A.J. and Bell, J.I.
                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates; 1 (bases 1 to 1143)
                                                                                   Screaton, G.R.
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                                                                                                                                                                                                                                                                             lymphocyte associated
                     Location/Qualifiers
1. .1143
/organism="Homo sapiens"
/db_xref="taxon:9606"
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/db_xref="G1:2071967"
/db_xref="G1:2071967"
/db_xref="G1:2071967"
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/db_xref="G1:2071967"
/translation="medepregcavanallizuligarangggtreproceduligary
/translation="medepregcysocysocysocyclocal-Hreffrilcsrrdp
Alencsavadtrogckempyecovsocysocyclocal-Hreffrilcsrrdp
CGFCLpgfrebebgdgcysocysocyclocal-Hreffrilcsrrdp
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CGFCLpgfrebebgdgcysocyclocal-Hreffrilcsrrdp
CGFCBAMMIOPGPOLYDVMDAVPARRWKEFVRTLGLREAE-EAVEYBIGRF
EDQQYEMLKRRRQQQPAGLGAVYAALERMGLDGCYEDLRSRLQRGP"
598. 599
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note="deletion compared to LARD 1a, deposited in GenBaccession Number U94501, probable skipping of putative exon 7 encoding the transmembrane domain" 378 c 378 g 199 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TNF-R1"
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/note="LARD-9; NGFR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /cell_type="lymphocyte"
1    .1143
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Best Local S
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                                                  TCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCCGCTGTGGCTGTAAGCCA 300
                                                                                             CCTTGCACGGAGCCCTGCGGCAACTCCACCTGCCTTGTGTGTCTCCCCAAGACACCTTCTTG
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GCCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCCTTCTACTGCCAA
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U94503
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Submitted (19-MAR-1997) Molecular Immunology Group, Institute of
Molecular Medicine, John Radcliffe Hospital, Headington, Oxford OX3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Screaton,G.R., Xu,X.N., Olsen,A.L., Cowper,A.E., Tan,R., McMichael,A.J. and Bell,J.I.
LARD: a new lymphoid-specific death domain containing receptor regulated by alternative pre-mRNA splicing
Proc. Natl. Acad. Sci. U.S.A. 94 (9), 4615-4619 (1997)
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Screaton, G.R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                /note="insertion compared to LARD la, deposited in GenBank Accession Number U94501, probable retained intron leading to premature transcriptional termination" 441 c 451 g 240 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /function="mediates apoptosis"
/note="LARD-2; similar to Fas and TNF-R1; possibl soluble; truncated before the transmembrane domai
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GEEGGNHPTPTSCFQCSGSRCSWLALWSPSCLGPP"
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LFCCRGCPAGHYLKAPCTEPGGNSTCLVCPQDTFLAWERNHNUSSCARCQACDEQASQV
ALENCSAVADTRCGGCRGMEYVEQVSSQVSSSSPYCQDCLDCGALHRHITRLLCSRADI
DCGTCLPGFYEHGDGCVSCPTSTLGSCPERCAAVCGWROSRWCAGNASGRTGMDRGEA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /product="lymphocyte associated receptor of
/protein_id="AAC51308.1"
/db_xref="GI:2071953"
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/db_xref="taxon:9606"
/cell_line="HeLa"
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Pred. No. 3.9e-129;
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                                       TCCCAGGTGGCGCTGGAGAACTGTTCAGCAGTGGCCGACACCCGCTGTGGCTGTAAGCCA,
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                                                                                GCCTGGGAGAACCACATAATTCTGAATGTGCCCGCTGCCAGGCCTGTGATGAGCAGGCC
                                                                                                                                     CTGGGGGCCCGGGCCCAGGGCACTCGTAGCCCCAGGTGTGACTGTGCCGGTGACTTC
GGCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCTTCTACTGCCAA
          GGCTGGTTTGTGGAGTGCCAGGTCAGCCAATGTGTCAGCAGTTCACCCCTTCTACTGCCAA
                                                                                                                           CCTTGCACGGAGCCCTGCGGCAACTCCACCTGCCTTGTGTGTCCCCCAAGACACCTTCTTG
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Sequence
AX331947
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Avalon Pharmaceuticals (US)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                        gene
                                                                                                                                                                                                                                                                                                                                                                                                  Cancer gene determination
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (12-NOV-1996) J. Kitson,
Research Centre, Gunnels Wood Road,
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nature 384 (6607), 97088617
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kitson, J., Raven, T., Jia
Pun, K.T., Grinham, C.J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  H.sapiens mRNA for WSL-LR, WSL-S1 and WSL-S2 proteins.
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1 (bases 1 to 1743)
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Mammalia; Eutheria; Primates; Catarrhini;
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                                                                                         /product="WSL-1R protein"
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ALENCSAVADTRCGCKPGWFVECQVSQCVSSPFYCQPCLDCGALHRHTRLLCSRRDT
DCGTCLRGFYEHGDGCVSCPFSTLGSCPERCAAVCGWRQMFWYQVLLAGLVVPLLLGA
TLTYTYRHCWPHKPLVTADEAGMEALTPPPATHLSPLDSAHTLLAPDDSSEKICTVQL
VGNSWTPGYPGTGEALCPQVTWSWDOLPSRALGPARAFTLSPESPAGSPAMMLQPGPQ
LYDVNDAVPARRWKEFVRTLGLREAEIEAVEVEIGLFRDQQYEMLKHWRQQQPAGLGA
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/gene="wsl-1"
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/db_xref="GI:1669692"
/db_xref="SWISS-PROT:Q930
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/tissue_type="lymphoid"
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/db_xref="taxon:9606"
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DCGTCLPGFYEHGDGCVSCPTSTLGSCPERCAAVCGWRQSRWCAGNARGRTGMDRGEA
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DCGTCLPGFYEHGDGCVSCPTVLGPGAPGWPCGPPPAWGHPDLHIPPLLASQAPGYCR
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